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<p>(21) International Application Number: <b>PCT/US96/09988</b></p> <p>(22) International Filing Date: <b>7 June 1996 (07.06.96)</b></p> <p>(30) Priority Data: <b>08/484,172 7 June 1995 (07.06.95) US</b></p> <p>(71) Applicant: MEDAREX, INC. [US/US]; 1545 Route 22 East, Annandale, NJ 08801-0992 (US).</p> <p>(72) Inventors: DEO, Yashwant, M.; 3009 Peacock Drive, Audubon, PA 19403 (US). GOLDSTEIN, Joel; 3804 Hana Road, Edison, NJ 08817 (US). GRAZIANO, Robert; 1046 County Road, Route 519, Frenchtown, NJ 08825 (US). SOMASUNDARAM, Chezian; Apartment 301, 1202 B. North Maxwell Street, Allentown, PA 18103 (US).</p> <p>(74) Agents: ARNOLD, Beth, E. et al.; Lahive &amp; Cockfield, 60 State Street, Boston, MA 02109 (US).</p>		<p>(81) Designated States: AL, AM, AT, AU, AZ, BB, BG, BR, BY, CA, CH, CN, CZ, DE, DK, EE, ES, FI, GB, GE, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, TJ, TM, TR, TT, UA, UG, UZ, VN, ARIPO patent (KE, LS, MW, SD, SZ, UG), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG).</p> <p><b>Published</b> <i>With international search report. Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.</i></p>	
<p>(54) Title: <b>THERAPEUTIC COMPOUNDS COMPRISED OF ANTI-Fc RECEPTOR ANTIBODIES</b></p> <p><b>b22 hinge region:</b></p> <p>[A]</p> <p>-ACT CAC ACA TGC CCA CCG TGC CCA — CH2 — CH3</p> <p>  T H T C P P C P</p> <p style="text-align: center;">↓</p> <p>[B]</p> <p>—<u>AntiII</u>—</p> <p>-ACT CAC ACA TGC CCA CCG TGA GGA TCC-</p> <p>  T H T C P P</p> <p style="text-align: center;">↓</p> <p>[C]</p> <p>—<u>Xhol</u>—    —<u>NdeI</u>—    —<u>SamIII</u>—</p> <p>-ACT CAC ACA TGC TCG AGC GTC CAC GCG GGC CGC TGA GGA TCC</p> <p>  T H T C S S L H G G R</p> <p>(57) Abstract</p> <p>Multispecific multivalent molecules which are specific to an Fc receptor (FcR), and therapeutic uses and methods for making the molecules are described.</p>			

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## Therapeutic Compounds Comprised of Anti-Fc Receptor Antibodies

### Background of the Invention

5        Immunoglobulins (Igs) are composed of two heavy and two light chains, each of which contains an NH<sub>2</sub>-terminal antigen-binding variable domain and a COOH-terminal constant domain responsible for the effector functions of antibodies. The COOH-terminal domains of Ig heavy chains form the Fc region and are involved in triggering cellular activities through interaction with specific receptors known as Fc receptors (FcRs). Fc  
10      receptors for all Ig classes, or isotypes, (e.g., IgG (Fc $\gamma$ R), IgE (Fc $\epsilon$ R), IgA (Fc $\alpha$ R), IgM (Fc $\mu$ R) and IgD (Fc $\delta$ R) have been identified. The different biological activities of antibodies of different isotypes are based in part on their ability to bind to different FcR expressed on different immune (effector) cells (Fridman, W.H. (Sept. 1991) The FASEB Journal Vol. 5. 2684-2690). Murine antibodies, which are directed against FcRs have been made (See e.g.  
15      U.S. Patent No. 4,954,617 entitled *Monoclonal Antibodies To Fc Receptors for Immunoglobulin G on Human Mononuclear Phagocytes* and International Patent Application Publication No. WO 91/05871 entitled *Monoclonal Antibody Specific For IgA Receptor*).

20      Murine monoclonal antibodies can be useful as human therapeutics and can be produced free of contamination by human pathogens such as the hepatitis or human immunodeficiency virus. However, use of murine monoclonal antibodies in some human therapies, have resulted in the development of an immune response to the "foreign" murine proteins. This response has been termed a human anti-mouse antibody or HAMA response (Schroff, R. et al. (1985), *Cancer Res.*, 45, 879-885) and is a condition which causes serum sickness in humans and results in rapid clearance of the murine antibodies from an individual's circulation. The immune response in humans has been shown to be against both the variable and the constant regions of murine immunoglobulins.  
25

30      Recombinant DNA technology can be used to alter antibodies, for example, by substituting specific immunoglobulin regions from one species with immunoglobulin regions from another species. Neuberger et al. (Patent Cooperation Treaty Patent Application No. PCT/GB85/00392) describes a process whereby the complementary heavy and light chain variable domains of an Ig molecule from one species may be combined with the complementary heavy and light chain Ig constant domains from another species. This process may be used to substitute the murine constant region domains to create a "chimeric" antibody which may be used for human therapy. A chimeric antibody produced as described by Neuberger et al. has a human Fc region for efficient stimulation of antibody mediated effector functions, such as complement fixation, but still has the potential to elicit an immune response in humans against the murine ("foreign") variable regions.  
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Winter (British Patent Application Number GB2188538A) describes a process for altering antibodies by substituting the complementarity determining regions (CDRs) with those from another species. This process may be used to substitute the CDRs from the 5 murine variable region domains of a monoclonal antibody with desirable binding properties (for instance to a human pathogen) into human heavy and light chain Ig variable region domains. These altered Ig variable regions may then be combined with human Ig constant regions to create antibodies which are totally human in composition except for the substituted murine CDRs. The "reshaped" or "humanized" antibodies described by Winter elicit a 10 considerably reduced immune response in humans compared to chimeric antibodies because of the considerably less murine components. Further, the half life of the altered antibodies in circulation should approach that of natural human antibodies. However, as stated by Winter, merely replacing the CDRs with complementary CDRs from another antibody which is 15 specific for an antigen such as a viral or bacterial protein, does not always result in an altered antibody which retains the desired binding capacity. In practice, some amino acids in the framework of the antibody variable region interact with the amino acid residues that make up the CDRs so that amino acid substitutions into the human Ig variable regions are likely to be required to restore antigen binding.

20 Bispecific molecules, (e.g., heteroantibodies) comprising an anti-Fc receptor portion and an anti-target portion have been formulated and used therapeutically, e.g., for treating cancer (e.g. breast or ovarian) or pathogenic infections (e.g., HIV) (See, e.g., International Patent Application Publication No. WO 91/05871 entitled *Bispecific Heteroantibodies With Dual Effector Functions*; and International Patent Application 25 Publication No. WO 91/00360 entitled *Bispecific Reagents for AIDS Therapy*). In addition, bispecific molecules, which recognize antigens and antigen presenting cells can be administered to a subject to stimulate an immune response (See, e.g., International Patent Application Publication No. WO 92/05793 entitled *Targeted Immunostimulation With Bispecific Reagents*).

Summary of the Invention

In one aspect, the invention features multispecific, multivalent molecules, which minimally comprise an anti-Fc receptor portion, an anti-target portion and optionally an anti-enhancement factor (anti-EF) portion. In preferred embodiments, the anti-Fc receptor portion is an antibody fragment (e.g., Fab or (Fab')<sub>2</sub> fragment), the anti-target portion is a ligand or antibody fragment and the anti-EF portion is an antibody directed against a surface protein involved in cytotoxic activity. In a particularly preferred embodiment, the recombinant anti-FcR antibodies, fragments or ligand are "humanized" (e.g., have at least a portion of a complementarity determining region (CDR) derived from a non-human antibody (e.g., murine) with the remaining portion(s) being human in origin).

In another aspect, the invention features methods for generating multispecific molecules. In one embodiment, both specificities are encoded in the same vector and are expressed and assembled in a host cell. In another embodiment, each specificity is generated recombinantly and the resulting proteins or peptides are conjugated to one another via sulfhydryl bonding of the C-terminus hinge regions of the heavy chain. In a particularly preferred embodiment, the hinge region is modified to contain only one sulfhydryl residue, prior to conjugation.

Recombinant antibodies and multispecific molecules generated therefrom can be engineered to have increased affinity and specificity. Further, humanized antibodies are typically less immunogenic when administered to a human. Other features and advantages of the present invention will become better understood by reference to the following Detailed Description and Claims.

Brief Description of the Drawings

Figure 1 is a diagram showing the nucleotide and amino acid residue sequences of a portion of the hinge region of a humanized Fcγ RI antibody, H22. [A] that was altered to produce a truncated single-sulfhydryl version [B] and then altered further to engineer two unique cloning sites [C]. Underlined nucleotides indicate changes from the previous sequence. Overlined nucleotides are the recognition sequences for the indicated restriction sites.

Figure 2 is a schematic representation of the heavy chain-EGF fusion expression construct pJG055.

Figure 3 is a schematic representation of the generation of anti-Fc receptor-ligand bispecific molecules.

5 Figure 4 is a schematic representation of the flow cytometric assay used for testing the activity of the humanized Fc  $\gamma$  receptor- epidermal growth factor fusion protein.

10 Figure 5 is a graph, which plots Mean Fluorescence Intensity (MFI) an indication of the binding of various concentrations of epidermal growth factor (EGF) fusion protein (H22-EGF fusion) and the fully humanized bispecific (BsAb) H447 to EGF receptor (EGFR) expressing 1483 cells.

15 Figure 6 is a graph, which plots the binding of various concentrations of the EGF fusion protein or the BsAb H447 to A431 cells in the presence and absence of murine antibody M425, which binds EGFR.

20 Figure 7 is a graph, which plots the antibody dependent cytotoxicity (ADCC) resulting from the binding of various concentrations of the EGF fusion protein, BsAb H447 or the H425 antibody to A431 cells.

25 Figure 8 is a bar graph which plots the ADCC resulting from the binding of EGF fusion protein, BsAb H447 or the H425 antibody in the presence of media alone, media containing 25% human serum (HS) or media containing a fab fragment of the Fc $\gamma$  receptor antibody m22.

30 Figure 9 is a schematic diagram representing the number of viable A431 cells cultured in the presence of various amounts of EGF, H22-EGF, the Fab fragment of H22 (H22 Fab), or the F(ab')<sub>2</sub> fragment of H425 (H425 F(ab')<sub>2</sub>).

Figure 10 shows the amino acid sequence of the H22Fd-HRG fusion protein.

35 Figure 11 is a histogram indicating the percentage of specific PC-3 or SKBr-3 tumor cell killing resulting from incubation of these cells with interferon- $\gamma$ -treated monocytes and a 1:3 or 1:30 dilution of supernatant from myeloma cells expressing an H22-heregulin fusion protein.

Figure 12 is a diagram indicating the percentage of PC-3 tumor cell lysis in the presence of monocytes and in the presence of various concentrations of H22-bombesin fusion protein concentrations.

Figure 13 is a schematic representation of the flow cytometric assay used for testing the activity of BsAb 447 generated either by the o-PDM or the DTNB method.

5 Figure 14 is a graph, which plots the MFI of various concentrations of o-PDM and DTNB derived BsAb 447 to EGFR and Fc<sub>Y</sub>RI expressing A431 cells.

Figure 15 is a graph, which plots the antibody dependent cytotoxicity resulting from the binding of o-PDM and DTNB derived BsAb 447 to A431 cells.

Figure 16 is a flow chart that depicts the construction of trispecific antibodies.

10 Figure 17 depicts the transformation of a bivalent, bispecific antibody into a trivalent, bispecific antibody. The bivalent, bispecific conjugate is reduced and mixed with o-PDM-treated 520C9 Fab' resulting in the TsAb.

Figure 18 depicts a bifunctional fluorescence-activated cell sorting assay for HER2/neu (panel A) and EGFR (panel B).

15 Figure 19 is a graph which plots the binding of various concentrations of antibody, either BsAb or TsAb, to target cells. Mean Fluorescence Intensity (MFI) increases as Ab binding increases. It shows that the TsAb bound both HER2/neu on SKBr-3 cells and soluble Fc<sub>Y</sub>RI simultaneously in a dose-dependent fashion.

20 Figure 20 is a graph that shows the TsAb bound both EGFR on A431 cells and soluble Fc<sub>Y</sub>RI simultaneously in a dose-dependent fashion. The assay is similar to that used in Figure 19.

25 Figure 21 is a graph that shows the TsAb, M22 x H425x 520C9, and the BsAb, M22 x 520C9 were capable of inducing ADCC of SKBR-3 cells but the BsAb, M22 x H425, was not. Various concentrations of antibodies were incubated with SKBR-3 cells and pre-activated PMNs.

Figure 22 is a graph that shows the TsAb, M22 x H425x 520C9, and the BsAb, M22 x H425 were capable of inducing ADCC of A431 cells but the BsAb, M22 x 520C9, was not. The assay was performed in a similar manner as the assay in Fig. 21.

30 Figure 23 is a flow chart for a whole blood modulation assay (panel A) and the results from the assay (panel B). This trivalent antibody rapidly modulates Fc<sub>Y</sub>RI from the surface of monocytes.

Figure 24, panel A, shows the amino acid sequence of oligonucleotides encoding the wildtype (TT830) and mutant (TT833) tetanus toxin peptides. Panel B is a diagram of an H22Fd-TT fusion protein.

5       Figure 25 panels A, B, and C represent flow cytometry analysis results showing binding of MDXH210, Fab22-TT830, and H22-TT833S to Fc $\gamma$ RI positive U937 cells, respectively. The dashed lines represent negative controls, the solid lines denote staining with the fusion proteins, and the dotted lines represent fusion protein binding blocked by murine mAb 22 F(ab')<sub>2</sub>.

10      Figure 26 is a schematic diagram showing the mean fluorescence intensity resulting from incubation of various amounts of the fusion proteins MDXH210, FAb22-TT830, and Fab22-TT833S to Fc $\gamma$ RI positive U937 cells.

15      Figure 27 is a graphic representation of the proliferation of T cells incubated with irradiated monocytes and various concentrations of TT830, Fab22-TT830, TT, or TT947, showing that the fusion protein Fab22-TT830 enhances the presentation of the Th epitope by about 1000 fold as compared to TT830.

20      Figure 28 represents a histogram showing the proliferation of T cells incubated with TT830 at 1000 nM or FAb22-TT830 at 10 nM and monocytes, preincubated or not with saturating amounts of mAb 22 F(ab')<sub>2</sub> prior to addition of the T cells and the antigen.

25      Figure 29 represents a histogram showing the proliferation of T cells incubated with monocytes and Fab22-TT830 at 5 nM or TT830 at 1000 nM in the absence (control) or presence of IgG.

30      Figure 30, panels A and B, are graphic representations showing the concentration of IFN- $\gamma$  (panel A) and IL-4 (panel B) in the supernatant of T cells cultured for 2 days with monocytes and various concentrations of TT830 or Fab22-TT830.

35      Figure 31 is a graphic representation depicting the proliferation of T cells incubated with monocytes and various concentrations of TT833S, Fab22-TT833S, or TT830.

Figure 32 is a graphic representation of the proliferation of T cells incubated for 2 days with TT830 and monocytes, preincubated overnight with various concentrations of TT833S.

Figure 33 is a graphic representation of the percent inhibition of proliferation of T cells incubated for 2 days with TT830 and monocytes, preincubated overnight with various concentrations of TT833S or FAb22-TT833S.

5                   Figure 34 is a histogram representing the proliferation of T cells incubated for 2 days with monocytes, which were first incubated with TT830 for 4 hours (Pre-pulse) and then incubated overnight with 10 $\mu$ M TT833S or 0.1 $\mu$ M Fab22-TT833S (Chase) prior to addition of the T cells.

10                  Figure 35 is a histogram representing the concentration of interferon- $\gamma$  (IFN- $\gamma$ ) and IL-4 in the supernatant of T cells cultured with monocytes and TT830, FAb22-TT830, TT833S, and Fab22-TT833S.

15                  Figure 36 is a graphic representation of the proliferation of T cells stimulated for one day with monocytes in medium alone, with TT833S, or with Fab22-TT833S and then restimulated with monocytes and various concentrations of TT830 for two days, indicating that TT833S and Fab22-TT833S do not lead to T cell anergy.

20                  Figure 37 is a graphic representation of two expression constructs encoding single chain bispecific molecules having one binding specificity for an Fc $\gamma$ RI (H22) and one binding specificity for a carcinoembryonic antigen (CEA) (constructs 321 and 323) and one expression construct encoding a single chain antibody having one binding specificity for an Fc $\gamma$ RI. The coding regions are under the control of the CMV promoter (CMV Pr). In addition to the variable regions from the heavy (VH) and light chains (VL) of the antibodies, 25 the proteins encoded by these constructs are fused to a peptide from c-myc (c-myc) and to a hexa-histidine peptide (H-6).

30                  Figure 38 shows a histogram indicating the level of binding of the single chain bispecific molecules H22-anti-CEA encoded by the expression constructs 321 (321-A5 and 321-B4) and 323 (323-B2 and 323-C4) and the single chain H22 antibody encoded by the construct 225 (225-C2) as measured by bispecific ELISA.

35                  Figure 39 shows the nucleic acid sequence of the single chain humanized anti-Fc $\gamma$ RI antibody and the amino acid sequence encoded by the nucleic acid.

                        Figure 40 shows the nucleic acid sequence of the single chain bispecific molecule having one binding specificity for the Fc $\gamma$ RI and one binding specificity for CEA and the amino acid sequence encoded by the nucleic acid.

Detailed Description*Multispecific molecules*

5 The instant invention relates to recombinant multispecific molecules. Multispecific molecules can include bispecific molecules comprised of an anti-Fc receptor portion and an anti-target portion, wherein at least one of said portions is constructed using recombinant DNA techniques. Multispecific molecules can also include molecules, which are comprised of more than one anti-Fc receptor portion or anti-target portion; or molecules  
10 10 comprised of at least one anti-Fc receptor, one anti-target portion and additionally a portion or portions that recognize another molecule, wherein at least one of said portions is constructed using recombinant DNA techniques.

An "anti-Fc receptor portion" refers to an antibody, a functional antibody  
15 fragment (e.g., Fab fragment) or a ligand that recognizes and binds an Fc receptor on an effector cell. Preferred antibodies for use in the subject invention bind the Fc receptor on an effector cell at a site which is not bound by endogenous immunoglobulin. Most preferably, the anti-Fc receptor portion binds a human Fc $\gamma$ R (i.e., Fc $\gamma$ RI, Fc $\gamma$ RII or Fc $\gamma$ RIII). Preferred humanized anti-Fc $\gamma$ R monoclonal antibodies are described in PCT application WO 94/10332  
20 and U.S. Patent No. 4,954,617, the teachings of which are fully incorporated herein by reference).

An "effector cell", as used herein refers to an immune cell. Specific effector cells express specific Fc receptors and carry out specific immune functions. For example,  
25 monocytes, macrophages, neutrophils and dendritic cells, which express Fc $\gamma$ RI are involved in both specific killing of target cells and presenting antigens to other components of the immune system. The expression of a particular FcR on an effector cell can be regulated by humoral factors such as cytokines. For example, expression of Fc $\gamma$ RI has been found to be up-regulated by interferon gamma (IFN- $\gamma$ ). This enhanced expression increases the cytotoxic  
30 activity of Fc $\gamma$ RI cells against targets.

The recombinant antibodies or antibody fragments, which specifically bind to an Fc receptor are preferably "humanized" i.e. derived from a human antibody, but having at least a portion of a complementarity determining region (CDR) derived from a non-human antibody. The portion being selected to provide specificity of the humanized antibody for a human Fc receptor. The humanized antibody has CDRs derived from a non-human antibody and the remaining portions of the antibody molecule are human.  
35

5        The antibody may be whole, i.e. having heavy and light chains or any fragment thereof, e.g., Fab or (Fab')<sub>2</sub> fragment. The antibody further may be a light chain or heavy chain dimer, or any minimal fragment thereof such as a Fv or a single chain construct as described in Ladner et al. (U.S. Patent No. 4,946,778, issued August 7, 1990), the contents of which is expressly incorporated by reference.

10      The humanized antibody or fragment may be any human antibody capable of retaining non-human CDRs. The preferred human antibody is derived from known proteins NEWM and KOL for heavy chain variable regions (VHs) and REI for Ig kappa chain, variable regions (VKs).

15      The portion of the non-human CDR inserted into the human antibody is selected to be sufficient for allowing binding of the humanized antibody to the Fc receptor. A sufficient portion may be selected by inserting a portion of the CDR into the human antibody and testing the binding capacity of the created humanized antibody using the enzyme linked immunosorbent assay (ELISA).

20      All of the CDRs of a particular human antibody may be replaced with at least a portion of a non-human CDR or only some of the CDRs may be replaced with non-human CDRs. It is only necessary to replace the number of CDRs required for binding of the humanized antibody to the Fc receptor. A non-human CDR derived from a murine monoclonal antibody (mab), mab 22, is described in International Patent Application Publication No. WO 94/10332, the contents of which are fully incorporated herein by reference. The mab 22 antibody is specific to the Fc receptor and further is described in U.S. 25 Patent No. 4,954,617, issued September 4, 1988, the contents of which are also expressly incorporated by reference. The humanized mab 22 antibody producing cell line was deposited at the American Type Culture Collection on November 4, 1992 under the designation HA022CL1 and has the accession no. CRL 11177.

30      An antibody can be humanized by any method, which is capable of replacing at least a portion of a CDR of a human antibody with a CDR derived from a non-human antibody. Winter describes a method which may be used to prepare the humanized antibodies of the present invention (UK Patent Application GB 2188638A, filed on March 26, 1987), the contents of which is expressly incorporated by reference. The human CDRs 35 may be replaced with non-human CDRs using oligonucleotide site-directed mutagenesis as described in International Patent Application Publication Number: WO 94/10332 entitled, *Humanized Antibodies to Fc Receptors for Immunoglobulin G on Human Mononuclear Phagocytes*.

In addition to an anti-Fc receptor portion, the claimed multispecific molecules can comprise an "anti-target portion", i.e. an antibody, a functional antibody fragment or a ligand that recognizes and binds a pathogen (e.g., virus, bacteria, fungi), a pathogen infected cell, a cancer or tumor cell (e.g., breast, ovarian, prostate, etc.) or other unwanted cell in a subject (e.g., a human or animal) or an antigen or modified form thereof. Additionally, the target portion may comprise or be directed against an antigen. A preferred embodiment contains an antigen that can be used to stimulate the immune system, for example, in instances of chronic infection, to deplete antigen in the circulation, and to treat tumors. A particularly preferred embodiment has an antigen that is attached to a multivalent molecule containing an anti-FcR antibody.

In a specific embodiment of the invention, the multispecific molecule contains a ligand. The ligand can be any ligand that interacts with a molecule. In a preferred embodiment, the ligand binds a protein, e.g., a surface protein on a target cell, such as a cancer cell. Preferred ligands include ligands to receptors, such as growth or differentiation factors. For example, a multivalent molecule can comprise an epidermal growth factor, or at least a portion or modified form that is capable of interacting with a receptor, e.g., an epidermal growth factor receptor. In another preferred embodiment of the invention, the ligand is a small peptide, such as bombesin, gastrin-releasing peptide (GRP), litorin, neuromedin B, or neuromedin C. The sequences of the peptides can be found, e.g., in U.S. Patent No. 5,217,955, the content of which is incorporated herein by reference. The ligand can also be a modified form of any of these peptides. The modification can increase binding to the receptor, decrease binding, or not affect the binding to a receptor. The modification of the ligand can also transform an agonist into an antagonist, such that the ligand inhibit rather than stimulate cell proliferation. The modification of the ligand can be an addition, a deletion, a substitution, or a modification of at least one amino acid.

In a specific embodiment of the invention, a multivalent or bispecific molecule comprises an antigen. As used herein, the term "antigen" means any natural or synthetic immunogenic substance, a fragment or portion of an immunogenic substance, a peptidic epitope, or a hapten.

In one embodiment of the invention, a bi- or multispecific molecule is employed to target an antigen to the cell to enhance the processes of internalization and presentation by these cells, and ultimately, to stimulate an immune response therein. In a specific embodiment, the bispecific binding agent specifically binds the antigen (either directly, to an epitope of the antigen, or indirectly, to an epitope attached to the antigen) and, at the same time, binds a surface receptor of an antigen-presenting cell which can internalize antigen for processing and presentation. In another embodiment, the antigen is linked to the

multi- or bispecific molecule and at the same time binds a surface receptor of an antigen-presenting cell. The receptor-binding component of these bi- or multispecific molecule (and thus the bi- or multispecific molecule, itself) binds the receptor of the antigen-presenting cell. In some instances, binding of the molecule occurs without the molecule substantially being 5 blocked by the natural ligand for the receptor. As a result, targeting of the antigen to the receptor will not be prevented by physiological levels of the ligand and the targeted receptor will remain capable of binding the ligand and functioning.

One type of antigen can be an allergen. An "allergen" refers to a substance 10 that can induce an allergic or asthmatic response in a susceptible subject. The list of allergens is enormous and can include pollens, insect venoms, animal dander dust, fungal spores and drugs (e.g. penicillin). Examples of natural, animal and plant allergens include proteins specific to the following genera: *Canine* (*Canis familiaris*); *Dermatophagoides* (e.g. *Dermatophagoides farinae*); *Felis* (*Felis domesticus*); *Ambrosia* (*Ambrosia artemisiifolia*; 15 *Lolium* (e.g. *Lolium perenne* or *Lolium multiflorum*); *Cryptomeria* (*Cryptomeria japonica*); *Alternaria* (*Alternaria alternata*); *Alder*; *Alnus* (*Alnus glutinosa*); *Betula* (*Betula verrucosa*); *Quercus* (*Quercus alba*); *Olea* (*Olea europaea*); *Artemisia* (*Artemisia vulgaris*); *Plantago* (e.g. *Plantago lanceolata*); *Parietaria* (e.g. *Parietaria officinalis* or *Parietaria judaica*); *Blattella* (e.g. *Blattella germanica*); *Apis* (e.g. *Apis mellifera*); *Cupressus* (e.g. *Cupressus 20 sempervirens*, *Cupressus arizonica* and *Cupressus macrocarpa*); *Juniperus* (e.g. *Juniperus sabina*, *Juniperus virginiana*, *Juniperus communis* and *Juniperus ashei*); *Thuya* (e.g. *Thuya orientalis*); *Chamaecyparis* (e.g. *Chamaecyparis obtusa*); *Periplaneta* (e.g. *Periplaneta americana*); *Agropyron* (e.g. *Agropyron repens*); *Secale* (e.g. *Secale cereale*); *Triticum* (e.g. *Triticum aestivum*); *Dactylis* (e.g. *Dactylis glomerata*); *Festuca* (e.g. *Festuca elatior*); *Poa* (e.g. *Poa pratensis* or *Poa compressa*); *Avena* (e.g. *Avena sativa*); *Holcus* (e.g. *Holcus lanatus*); *Anthoxanthum* (e.g. *Anthoxanthum odoratum*); *Arrhenatherum* (e.g. *Arrhenatherum elatius*); *Agrostis* (e.g. *Agrostis alba*); *Phleum* (e.g. *Phleum pratense*); *Phalaris* (e.g. *Phalaris arundinacea*); *Paspalum* (e.g. *Paspalum notatum*); *Sorghum* (e.g. *Sorghum halepensis*); and *Bromus* (e.g. *Bromus inermis*). 25

30 Many allergens are found in airborne pollens of ragweed, grasses, or trees, or in fungi, animals, house dust, or foods. As a class, they are relatively resistant to proteolytic digestion. Preferable allergens are those which bind to IgE on mast cells and basophils, thereby causing a type I anaphylaxis hypersensitivity reaction. When at least one specificity 35 of the multivalent agent is for an epitope of the high affinity Fc receptor that is outside the ligand binding domain for IgG, this bispecific binding agent can decrease hypersensitivity in a subject. This is accomplished when the bispecific binding agent competes for an IgE-binding allergen before the allergen binds to IgE on a mast cell or basophil, thereby reducing the possibility of a type I hypersensitivity reaction. In addition, as a result of directing

allergen to Fc<sub>Y</sub>R, a state of T cell tolerance to the allergen may be induced which interferes with IgE-mediated type I reactions. Tolerance can be accomplished by inducing IgG which competes with IgE for binding to allergen using doses of allergen substantially lower than those currently used.

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In some cases, it may be desirable to couple a substance which is weakly antigenic or nonantigenic in its own right (such as a haptene) to a carrier molecule, such as a large immunogenic protein (e.g., a bacterial toxin) for administration. In these instances, the bispecific binding reagent can be made to bind an epitope of the carrier to which the 10 substance is coupled, rather than an epitope of the substance itself.

The antigen that can be linked either directly, or indirectly, to a multi- or bispecific molecule of the invention can be soluble or particulate; it may carry B cell epitopes, T cell epitopes or both. The antigen can be bacterial, viral or parasitic in origin.

15 Often, the antigen will comprise a component of the surface structure of a pathogenic organism. For example, the antigen can comprise a viral surface structure such as an envelope glycoprotein of human immunodeficiency virus (HIV) or the surface antigen of hepatitis virus. In addition, the antigen can be associated with a diseased cell, such as a tumor cell, against which an immune response may be raised for treatment of the disease.

20 The antigen can comprise a tumor-specific or tumor-associated antigen, such as the Her-2/new proto-oncogene product which is expressed on human breast and ovarian cancer cells (Slamon et al. (1989) Science 244:707).

25 The cells of a subject can be exposed *in vitro* or *in vivo* to the multivalent molecules of the invention. The multivalent molecule can be used to target an antigen to antigen-presenting cells in culture. Immunocompetent cells are separated and purified from patient blood. The cells are then exposed to a multivalent molecule comprising the antigen or the cells can be exposed to the antigen together with a multivalent molecule having a binding specificity for the antigen. Targeted antigen-presenting cells will process the antigen and 30 present fragments on their surface. After stimulation, the cells can be returned to the patient.

35 The method of this invention can be used to enhance or reinforce the immune response to an antigen. For example, the method is valuable for the treatment of chronic infections, such as hepatitis and AIDS, where the unaided immune system is unable to overcome the infection. It can also be used in the treatment of the acute stages of infection when reinforcement of immune response against the invading organism may be necessary.

The method can be used to reduce the dose of antigen required to obtain a protective or therapeutic immune response or in instances when the host does not respond or

responds minimally to the antigen. Although generally desirable, the lowering of effective dose can be especially desirable when the antigen is toxic to the host such as is the case for allergies. Methods and uses for using bi- or multispecific molecules comprising an antigen or comprising an ligand, e.g., an antibody interacting with an antigen, are further described in the published PCT application PCT/US91/07283.

In another embodiment of the invention, a multispecific molecule comprises an antigen that has been modified, such that its effect on T cell activation is modified upon presentation of the modified antigen to the T cell by an antigen presenting cell. Allan et al. have in fact shown that substitution of one or more amino acids of a peptide that stimulates T cells, e.g., stimulates T cell proliferation, can result in an antigen which fails to stimulate the T cell or which induces anergy in the T cell. Such modified peptides are termed Altered Peptide Ligands (APL). Accordingly, such APLs can be linked to bispecific or multispecific molecules having at least one binding specificity for the Fc<sub>Y</sub>RI. Upon phagocytosis of these molecules by antigen presenting cells and presentation to T cells, the proliferation of the T cells may be inhibited or anergized. Accordingly, administration to a subject of a multispecific molecule comprising (a) at least one altered peptide of an antigen which normally stimulates T cells, but which upon modification induces anergy of the T cells, and (b) at least one anti-Fc<sub>Y</sub>RI antibody will result in induction of tolerance of the subject to the antigen. Thus, such multi- or bispecific molecules can be used to tolerize a subject to a variety of antigens, e.g., auto-antigens. Thus, depending on the antigen used, the methods of the invention provide methods for increasing an immune response, i.e., by using an antigen which stimulates T cells, and the invention also provides methods for reducing an immune response, either by inhibiting T cell stimulation or by inducing anergy of the T cells.

The multispecific, multivalent molecules of the invention may also include an "anti-enhancement factor (anti-EF) portion". The "anti-enhancement factor portion" can be an antibody, functional antibody fragment or a ligand that binds to an antigen and thereby results in an enhancement of the effect of the anti-Fc receptor portion or the anti-target portion. The "anti-enhancement factor portion" can bind an Fc receptor or a target. A multivalent molecule comprising an anti-target portion that binds to one target cell antigen and an anti-enhancement factor portion that binds to a different target antigen is particularly useful where the target cell undergoes antigen modulation or antigenic variation (e.g., as has been described for certain parasites (such as trypanosomes). Alternatively, the anti-enhancement factor portion can bind an entity that is different from the entity to which the anti-target or anti-Fc receptor portion binds. For example, the anti-enhancement factor portion can bind a cytotoxic T-cell (e.g. via CD2, CD3, CD8, CD28, CD4, CD40, ICAM-1 or other immune cell that results in an increased immune response against the target).

Methods for Making Multispecific Molecules

The multispecific molecules described above can be made by a number of methods. For example, both specificities can be encoded in the same vector and expressed and assembled in the same host cell. This method is particularly useful where the multispecific molecule is a ligand x fab fusion protein as described in the following Example 2. A 5 bispecific molecule of the invention can also be a single chain bispecific molecule, such as a single chain bispecific antibody, a single chain bispecific molecule comprising one single chain antibody and a ligand, or a single chain bispecific molecule comprising two ligands.

10 Multivalent molecules can also be single chain molecules or may comprise at least two single chain molecules. Methods for preparing bi- or multivalent antibodies are for example described in U.S. Patent Number 5,260,203; U.S. Patent Number 5,455,030; U.S. Patent Number 4,881,175; U.S. Patent Number 5,132,405; U.S. Patent Number 5,091,513; U.S. Patent Number 5,476,786; U.S. Patent Number 5,013,653; U.S. Patent Number 5,258,498; 15 and U.S. Patent Number 5,482,858.

Binding of the single chain molecules to their specific targets can be confirmed by bispecific ELISA as described in the Examples herein.

20 Alternatively, each specificity of a multispecific molecule can be generated separately and the resulting proteins or peptides conjugated to one another. For example, two humanized antibodies can be conjugated via sulphydryl bonding of the C-terminus hinge regions of the two heavy chains. In a particularly preferred embodiment, the hinge region is modified to contain an odd number of sulphydryl residues, preferably one, prior to 25 conjugation.

The bispecific molecules of the present invention can be prepared by conjugating the anti-FcR and anti-target portions using methods described in the following Example or those well-known in the art. For example, a variety of coupling or cross-linking 30 agents can be used for covalent conjugation. Examples of cross-linking agents include protein A, carbodiimide, N-succinimidyl-S-acetyl-thioacetate (SATA), N-succinimidyl-3-(2-pyridyldithio)propionate (SPDP), and sulfosuccinimidyl 4-(N-maleimidomethyl) cyclohexane-1-carboxylate (sulfo-SMCC) (see e.g., Karpovsky et al. (1984) J. Exp. Med. 160:1686; Liu, MA et al. (1985) Proc. Natl. Acad. Sci. USA 82:8648). Other methods 35 include those described by Paulus (Behring Ins. Mitt. (1985) No. 78, 118-132); Brennan et al. (Science (1985) 229:81-83), and Glennie et al. (J. Immunol. (1987) 139: 2367-2375). Preferred conjugating agents are SATA and sulfo-SMCC, both available from Pierce Chemical Co. (Rockford, IL).

Therapeutic Uses for Multispecific Molecules

Based on their ability to bind FcR bearing immune cells and specific target cells, a specific multispecific molecule can be administered to a subject to treat or prevent a variety of diseases or conditions, including: cancer (e.g., breast, ovarian, small cell carcinoma of the lung), pathogenic infections (e.g., viral (such as HIV)), protozoan (such as Toxoplasma gondii), fungal (such as candidiasis); an autoimmunity (e.g. immune thrombocytopenia purpura and systemic lupus). The multispecific multivalent can also be administered prophylactically to vaccinate a subject against infection by a target cell.

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For use in therapy, an effective amount of an appropriate multispecific molecule can be administered to a subject by any mode that allows the molecules to exert their intended therapeutic effect. Preferred routes of administration include oral and transdermal (e.g., via a patch). Examples of other routes of administration include injection (subcutaneous, intravenous, parenteral, intraperitoneal, intrathecal, etc.). The injection can be in a bolus or a continuous infusion.

A multispecific molecule can be administered in conjunction with a pharmaceutically acceptable carrier. As used herein, the phrase "pharmaceutically acceptable carrier" is intended to include substances that can be coadministered with a multispecific molecule and allows the molecule to perform its intended function. Examples of such carriers include solutions, solvents, dispersion media, delay agents, emulsions and the like. The use of such media for pharmaceutically active substances are well known in the art. Any other conventional carrier suitable for use with the molecules falls within the scope of the instant invention.

The language "effective amount" of a multispecific molecules refers to that amount necessary or sufficient to realize a desired biologic effect. For example, an effective amount of a multispecific molecule, in which the anti-target portion recognizes a pathogenic cell could be that amount necessary to eliminate a tumor, cancer, or bacterial, viral or fungal infection. The effective amount for any particular application can vary depending on such factors as the disease or condition being treated, the particular multispecific molecule being administered, the size of the subject, or the severity of the disease or condition. One of ordinary skill in the art can empirically determine the effective amount of a particular multispecific molecule without necessitating undue experimentation.

The following invention is further illustrated by the following examples, which should not be construed as further limiting. The contents of all references, pending

patent applications and published patents, cited throughout this application are hereby expressly incorporated by reference.

### Examples

5

**Example 1: Production of Bispecific Antibody Comprising Murine or Humanized Antibodies Specific for an Fc Receptor and an Anti-her 2 neu Antibody**

10

#### Monoclonal Antibodies

The anti-Fc<sub>Y</sub>RI monoclonal antibodies (mAbs), M22, M32.2 and 197 were purified from hybridoma supernatant by ion exchange chromatography and DZ33, a human anti-HIV-1 IgG1 mAb, was purified from hybridoma supernatant by protein A affinity chromatography (Pharmacia, Piscataway, NJ) and gel filtration. M32.2 was deposited at the 15 American Type Culture Collection, 12301 Parklawn Drive, Rockville, MD 20852, on July 1, 1987 and has been designated with ATCC Accession No. HB9469.

20

#### Cell Lines

The murine myeloma NSO (ECACC 85110503) is a non-Ig synthesizing line 25 and was used for the expression of recombinant mAbs. NSO cells were cultivated in DMEM plus 10% fetal bovine serum (FBS, Gibco, Paisley, U.K.). SKBR-3 is a human breast carcinoma cell line which overexpresses the HER2/neu protooncogene (ATCC, Rockville, MD) and was cultivated in Iscove's Modified Dulbecco's Medium (IMDM, Gibco, Grand Island, NY). U937 is a monocytoid cell line that expresses Fc<sub>Y</sub>RI and was obtained from ATCC and grown in RPM-1640 plus 10% FBS (Gibco, Grand Island, NY).

30

#### Cloning Murine Immunoglobulin V Region Genes

Cytoplasmic RNA from the murine hybridoma 22 was prepared as described in Favaloro et al. (Favaloro, J., R. Treisman and R. Kamen (1982) *Transcription maps of polyoma-specific RNA: analysis by two-dimensional S1 gel mapping*. Meth. Enzymol. 65:718). The Ig V region cDNAs were made from RNA via reverse transcription initiated from primers CG1FOR and CK2FOR as described in International Patent Application Publication Number WO 94/10332 entitled, *Humanized Antibodies to Fc Receptors for Immunoglobulin G on Human Mononuclear Phagocytes*. The cDNA synthesis was 35 performed under standard conditions using 100 U MMLV reverse transcriptase (Life Technologies, Paisley, UK). The V<sub>H</sub> and V<sub>K</sub> cDNAs were amplified by PCR, (Orlandi, R., D.H. Güssow, P.T. Jones and G. Winter (1989) (Cloning immunoglobulin variable domains for expression by the polymerase chain reaction), Proc. Natl. Acad. Sci. USA 86:3833), using the cDNA primers in concert with SH2BACK and VK7BACK as described in International

Patent Application Publication Number WO 94/10332. Amplified  $V_H$  and  $V_K$  DNA were purified, cloned into M13, and sequenced by the dideoxy method using T7 DNA polymerase (Pharmacia, Piscataway, NJ).

5

Construction of Chimeric Antibody Genes

To facilitate cloning of murine V region DNA into expression vectors, restriction sites were placed in close proximity to the termini of both M22 V region genes. For  $V_H$ , a 5' *Pst*I site and a 3' *Bst*EII site were introduced into a cloned murine  $V_H$  gene by PCR using VH1BACK and VH1FOR (Id.). For  $V_K$  a 5' *Pvu*II site and a 3' *Bgl* II site were 10 introduced into a cloned murine  $V_K$  gene by PCR using primers VK1BACK and VK1FOR (Id.). In some instances, these primers changed one or more amino acids from those naturally occurring. These V region genes (ChVH and ChVK) were cut with the appropriate restriction enzymes and cloned into M13VHPCR1 and M13VKPCR1 (Id.) which contain an Ig promoter, signal sequence and splice sites. The DNA were excised from M13 as *Hind*III- 15 *Bam*HI fragments and cloned into the expression vectors pSVgpt and pSVhyg containing human IgG1, (Takahashi, N. et al., (1982), Structure of human immunoglobulin gamma genes: implications for evolution of a gene family, *Cell*, 29:671), and human kappa constant, (Hieter, R.A. et al., (1980) Cloned human and mouse kappa immunoglobulin constant and J region genes conserve homology in functional segments, *Cell* 22:197), region genomic DNA.

20

Construction of Humanized Antibody Genes

Two humanized heavy chains were constructed and were based on human  $V_H$ s of NEWM, (Poljak, R.J. et al., Amino acid sequence of the  $V_H$  region of a human myeloma immunoglobulin, (IgG New), *Biochemistry*, 16:3412), and KOL,(Marquat, M. et al., (1980) Crystallographic refinement and atomic models of the intact immunoglobulin molecule Kol and its antigen-binding fragment at 3.0A and 1.9A resolution, *J. Mol. Biol.* 141:369. The humanized light chain was derived from the human Bence-Jones protein REI, (Epp, O. et al, (1974) Crystal and molecular structure of a dimer composed of the variable portion of the Bence-Jones protein REI, *Eur. J. Biochem.* 45:513), with some framework 25 region (FR) changes. The modifications were made to make the  $V_K$  domain more typical of human subgroup I, and included replacement of Thr39, Leu104, Gln105 and Thr107 with Lys39, Val104, Glu105 and Lys107. In addition, Met4 was changed to Leu4 to 30 accommodate a *Pvu*II restriction site.

35

DNA containing the NEWM  $V_H$  and REI  $V_K$  FRs with irrelevant CDRs were cloned into the vectors M13VHPCR1 and M13VKPCR1 (Favaloro et al. *Supra*). DNA encoding the KOL  $V_H$  was constructed by a series of sequential PCRs, using oligodeoxyribonucleotides encoding KOL FR amino acids and irrelevant CDRs. The constructs were then cloned into M13VHPCR1.

Oligodeoxyribonucleotides were synthesized to encode the mAB M22 CDRs which were flanked by nucleotides corresponding to the human FRs. For the humanized  $V_H$  based on NEWM, the primers included murine FR amino acids Phe27, Ile28 and Arg71 since 5 these were likely to influence antigen binding, (Chothia, C. and A.M. Lesk (1987), Canonical structures for the hypervariable regions of immunoglobulins, *J. Mol. Biol.*, **196**:901; Tramontano, A. et al., (1990), Framework residue 71 is a major determinant of the position and conformation of the second hypervariable region in  $V_H$  domains of immunoglobulins, *J. Mol. Biol.*, **215**:175). For the humanized  $V_{\kappa}$ , murine amino acid Phe71 was similarly 10 included as a residue capable of affecting affinity, (Foote, J. and G. Winter, (1992), Antibody framework residues affecting the conformation of the hypervariable loops, *J. Mol. Biol.*, **224**:487. No murine FR residues were included in the KOL  $V_H$ . Oligodeoxyribonucleotides were 5' - phosphorylated and with the M13 universal forward primer annealed to the human 15 V region genes cloned in M13 in reactions containing M13 ssDNA template. The DNA was extended and ligated with 2.5 U T7 DNA polymerase (United States Biochemicals, Cleveland, OH) and 0.5 U T4 DNA ligase (Gibco BRL, Grand Island, NY). The mutated strand was preferentially amplified from the extension/ligation mixture using M13 reverse sequencing primer with 1 U Vent DNA polymerase (New England Biolabs, Beverly, MA) and was then amplified by PCR using both M13 forward and reverse primers. Product DNA 20 was cut with *Bam*H1 and *Hind*III, cloned into M13 and triple CDR-grafted mutants identified by DNA sequencing.

M13 clones containing the humanized V regions were sequenced in their 25 entirety to ensure the absence of spurious mutations. RF DNA from the confirmed clones was digested with *Hind*III and *Bam*H1, cloned into pSVgpt or pSVhyg and human IgG1 or human kappa constant regions added exactly as described for the construction of the chimeric antibody genes.

#### Expression and Purification of Recombinant mAbs

30 Heavy (5  $\mu$ g) and light (10  $\mu$ g) chain expression vectors were digested with *Pvu*I, ethanol precipitated and dissolved in 50  $\mu$ l water. NSO cells ( $1-2 \times 10^7$ ) were harvested by centrifugation, resuspended in 0.5 ml DMEM and mixed with the DNA in a 0.4 cm electroporation cuvette. After 5 min. on ice the cells were given a single pulse of 170 V, 960  $\mu$ F (GenePulser, Bio-Rad, Melville, NY) and incubated further for 15 min. on ice. The 35 cells were allowed to recover in DMEM for 24-48 hours. The medium was then made selective by the addition of mycophenolic acid (0.8  $\mu$ g/ml) and xanthine (250  $\mu$ g/ml). Aliquots of 200  $\mu$ l were distributed into 96-well plates. After a further 10-12 days, cells from the wells containing the highest levels of antibody measured by ELISA were selected and cloned by limiting dilution.

Antibodies were purified from overgrown cultures by protein A affinity chromatography (Boehringer Mannheim, Lewes, U.K.) Concentrations were determined by measuring  $A_{280\text{nm}}$  and confirmed by ELISA and SDS-PAGE.

5

ELISA for Measurement of Antibody Binding

The wells of a microtiter plate were coated with goat anti-human IgM antibodies (Sera-Lab, Crawley Down, U.K.) in 50 mM bicarbonate buffer, pH 9.6. The plate was blocked with 1% BSA and followed by the addition of a soluble fusion protein consisting of the extracellular domain of human Fc $\gamma$ RI and human IgM heavy chain (sFc $\gamma$ RI- $\mu$ ) obtained from transiently transfected COS cells (the expression vector was kindly provided by Dr. Brian Seed, Massachusetts General Hospital, Boston, MA). Recombinant 22 or control mAbs were then added in the presence of excess (2.2  $\mu\text{g}/\text{well}$ ) human IgG1 antibodies (Sigma, St. Louis, MO) that contained  $\lambda$  light chains to block the non-specific binding of the test mAbs via their Fc portion. Bound 22 mAbs were detected with peroxidase-labeled goat anti-human kappa chain antibodies (Sera-Lab, Crawley Down, U.K.) and o-phenylenediamine.

20

Fluoresceination of Antibodies

The pH of mAb solution was adjusted to 9.3 by the addition of 0.1M  $\text{Na}_2\text{CO}_3$ . Fluorescein iso-thiocyanate (FITC) (Sigma, St. Louis, MO) was dissolved in DMSO at a concentration of 2mg/ml. Forty  $\mu\text{g}$  of FITC was added for each milligram of mAb and incubated for two hours at room temperature. The fluoresceinated mAb was separated from the free FITC by G-25 chromatography.

25

Preparation of Blood Cells

Buffy coats were prepared from heparinized whole venous blood. Whole blood was diluted with RPMI containing 5% dextran at a ratio of 2.5:1 (v/v). The erythrocytes were allowed to sediment for 45 minutes on ice, then the cells in the supernatant were transferred to a new tube and pelleted by centrifugation. The residual erythrocytes were removed by hypotonic lysis. The remaining lymphocytes, monocytes and neutrophils were kept on ice until use in binding assays. For some experiments, neutrophils were separated from mononuclear cells by ficoll hypaque (Pharmacia, Piscataway, NJ) gradient separation. To up-regulate Fc $\gamma$ RI, neutrophils and mononuclear cells were treated with cytokines. Cultures of mononuclear cells were incubated at 37°C, 5%  $\text{CO}_2$  for 48 hours in teflon dishes at  $4 \times 10^6$  cells/ml of RPMI containing 2.5% normal human serum type AB (Sigma, St. Louis, MO) and 500 IRU/ml IFN- $\gamma$  (R&D Systems, Minneapolis, MN). Neutrophils were cultured for 48 hours (37°C, 5%  $\text{CO}_2$ ) in AIM V media (Gibco, Grand Island, NY) with 50 ng/ml G-CSF (Kindly provided by R. Repp, U. of Erlanger, Germany) and 500 IRU/ml

IFN- $\gamma$ .

Flow Cytometry

Cell binding assays were performed using 96-well microtiter plates as previously described, (Guyre, P.M. et al., Monoclonal antibodies that bind to distinct epitopes on Fc $\gamma$ R are able to trigger receptor function. *J. Immunol.*, **143**:1650). Briefly, cells were washed in PBS, pH 7.4 containing 2mg/ml BSA and 0.05% Na<sub>3</sub>N (PBA), and adjusted to 2.0 x 10<sup>7</sup> cells/ml with PBA. FITC-labeled and unconjugated antibodies were prepared in PBA. Cells (25  $\mu$ l), antibody (25  $\mu$ l) and human serum (25  $\mu$ l), or human IgG (10 mg/ml, Sigma, St. Louis, MO) (25  $\mu$ l), or PBA (25  $\mu$ l) were added to the microtiter plate, and left on ice for 45-60 minutes. Unbound antibody was removed from the wells by washing the cells 3 times with PBA. The cells were fixed with 1% paraformaldehyde. Cell associated fluorescence was analyzed on a Becton Dickinson FACScan.

15 BsAb Coupling Procedure

BsAb were constructed using the method of Glennie et al, (Glennie, M.J. et al., (1987), Preparation and performance of bispecific F(ab' gamma)<sup>2</sup>, antibody containing thioether-linked Fab' gamma fragments, *J. Immunol.*, **139**:2367). mAbs 22 (both murine and humanized) and 520C9 (anti-HER2/neu) antibodies were produced by *in vitro* cultivation of the respective hybridoma cells. The antibodies were separately digested with pepsin to F(ab')<sub>2</sub>, and subsequently reduced to Fab' by addition of 10 mM mercaptoethanolamine (MEA) for 30 minutes at 30°C. The Fab' fragments were applied to a Sephadex G-25 column equilibrated in 50mM Na Acetate, 0.5mM EDTA, pH 5.3 (4°C). Ortho-phenylenedimaleimide (o-PDM, 12mM) dissolved in dimethyl formamide and chilled in a 20 methanol/ice bath was added (one half volume) to the murine 22 Fab' in the case of M 22 x 520C9, and to 520C9 Fab' in the case of H 22 x 520C9 and incubated for 30 minutes on ice. The Fab'-maleimide was then separated from free o-PDM on Sephadex G-25 equilibrated in 50mM Na Acetate, 0.5mM EDTA, pH 5.3 (4°C). For preparation of the BsAbs, the M22 25 Fab'-maleimide was added to the 520C9 Fab' or the 520C9 Fab'-maleimide was added to H22 Fab' at a 1:1 molar ratio. The reactants were concentrated under nitrogen to the starting volume using a Diaflo membrane in an Amicon chamber (all at 4°C). After 18 hours the pH was adjusted to 8.0 with 1M Tris-HCl, pH 8.0. The mixture was then reduced with 10mM MEA (30 minutes, 30°C) and alkylated with 25 mM iodoacetamide. The bispecific F(ab')<sub>2</sub> 30 was separated from unreacted Fab's and other products by a Superdex 200 (Pharmacia, Piscataway, NJ) column equilibrated in PBS.

Antibody Dependent Cellular Cytotoxicity (ADCC)

The HER2/neu over-expressing human breast carcinoma cells, SKBR-3, were used as targets for lysis by cytokine activated neutrophils (see preparation of blood cells).

Targets were labeled with 100  $\mu$ Ci of  $^{51}\text{Cr}$  for 1 hour prior to combining with neutrophils and antibodies in a U-bottom microtiter plate. After incubation for 5 hours at 37°C supernatants were collected and analyzed for radioactivity. Cytotoxicity was calculated by the formula: % lysis = (experimental CPM - target leak CPM/detergent lysis CPM - target leak CPM) x 100%. Specific lysis = % lysis with antibody - % lysis without antibody. Assays were performed in triplicate.

#### Superoxide Induction

U937 cells were used for measuring the ability of H22 to trigger a superoxide burst via Fc $\gamma$ RI, (Pfefferkorn, L.C. and G.R. Yeaman (1994), Association of IgA-Fc receptors (Fc x R) with Fc $\epsilon$  RI $\gamma$  2 subunits in U937 cells, *J. Immunol.* **153**:3228; Hallet, H.B. and A.K. Campbell (1983). Two distinct mechanisms for stimulating of oxygen - radical production in polymorphonuclear leucocytes, *Biochem J.* **216**:459). U937 cells were cultured for five days in RPMI-1640 (Gibco, Grand Island, NY) with 10% FBS (Hyclone, Logan, UT) in the presence of 100 U/ml IFN- $\gamma$  (Genentech, S. San Francisco, CA) to induce differentiation and increased expression of Fc $\gamma$ RI. On the day of the experiment, these differentiated cells were incubated for 20 minutes in fresh RPMI-1640 with 10% FBS at 37°C. The cells were then pelleted and resuspended at a concentration of  $3 \times 10^6$  cells/ml in PBS supplemented with 1mM CaCl<sub>2</sub>, 1mM MgCl<sub>2</sub>, 11mM glucose, and 100 $\mu$ g/ml BSA (Sigma, St. Louis, MO). To trigger the release of superoxide, 100 $\mu$ l of cells were added to 100 $\mu$ l of a reaction solution containing 0.1mM luminol (Sigma, St. Louis, MO), 0.5mM sodium vanadate (Sigma, St. Louis, MO), and either mAb M22, H22, or 197 and placed in the luminometer at 22° C. Measurements of the spontaneous production of superoxide were made every 30 to 40 seconds starting immediately following the addition of the cells to the reaction solution in the luminometer. To compare the superoxide triggered by crosslinking Fc $\gamma$ RI with M22, H22 or 197, each mAb was used at a concentration of 10 $\mu$ g/ml. The production of superoxide in mV/sec was monitored for 20 minutes. MAb M22, M32.2 and 197 were added at various concentrations to establish the dose-responsiveness of superoxide production.

30 Results

#### Murine Ig V Region Genes

Ig V region cDNAs were prepared from M22 hybridoma RNA using primers specific for murine heavy and kappa constant regions and were amplified by PCR with the additional use of a series of primers based on sequences of known signal and/or 5' sequences of mature V regions. PCR products of the expected sizes for V<sub>H</sub> and V<sub>K</sub> were obtained using the SH2BACK/CG1FOR and VK7BACK/CK2FOR primer combinations. Amplified DNA was digested with appropriate restriction enzymes, cloned into M13 and the sequence in both directions determined from at least 24 independent clones. The deduced amino acid

sequences are shown in SEQ. ID Nos. 29 and 30. The 4 N-terminal residues of  $V_K$  are encoded by the VKBACK primer.

The M22  $V_H$  and  $V_K$  are members of murine heavy chain subgroup IIID and 5 kappa subgroup I, (Kabat, E.A. et al., (1991), Sequences of Proteins of Immunological Interest, 5th Ed., U.S. Department of Health and Human Services), respectively. Apart from the residue at L97, the amino acid sequence of the M22  $V_K$  is identical to that from the murine anti-IgG mAb A17 (Shlomchik, M. et al., Variable region sequences of murine IgM anti-IgG monoclonal autoantibodies (rheumatoid factors). II Comparison of hybridomas 10 derived bylipopolysaccharide stimulation and secondary protein immunization, J. Exp. Med. 165:970).

**Humanized mAbs and Initial Characterization of their Binding**

M22  $V_H$  FR showed greater homology (79%) to KOL (human subgroup III) 15 than to NEWM (57%) (human subgroup II). To see how this difference might affect binding, heavy chains were constructed based either on NEWM  $V_H$  including the murine residues Phe27, Ile28 and Arg71, or on KOL  $V_H$  with no murine FR amino acids. Both humanized  $V_H$  were partnered with the same REI-derived humanized light chain.

20 The affinity of the humanized mAbs was initially assessed by ELISA measuring the binding to Fc $\gamma$ RI/IgM heavy chain fusion protein. The data showed that the KOL  $V_H$ /REI  $V_K$  mAb had the same binding as the chimeric mAb whereas the NEWM  $V_H$ /REI  $V_K$  mAb exhibited an approximate 5- fold lower affinity. The low binding of a nonspecific human IgG1 mAb showed that >95% of binding of the humanized mAbs was via 25 the Fv portion rather than through the Fc domain.

30 While additional changes to the NEWM FR would be expected to restore binding affinity these could create novel epitopes which might provoke an unwanted immunological response. The KOL  $V_H$ /REI  $V_K$  mAb, designated H22, was therefore chosen for further examination of its binding characteristics.

**Functional Characterization of mAbH22**

A series of binding experiments were performed to establish the specificity and isotype of the H22 antibody. Peripheral blood leukocytes stained with fluorescein-conjugated M22 or H22 demonstrated specific binding to monocytes with approximately  $10^4$  35 binding sites per cell. In contrast, lymphocytes or unstimulated neutrophils had little or no specific binding (Table 1):

Table 1: Specific Binding of H22 to Monocytes

Antibody	Monocytes	Lymphocytes	PMNs
M22	10,000 <sup>a</sup>	<1000	<1000
H22	10,500	<1000	<1000

<sup>a</sup>Antibody sites per cell, average of duplicates

5 To demonstrate that the H22 binds to Fc<sub>Y</sub>RI at the same site as M22 and that it also binds as a ligand at the Fc binding domain, competition experiments with two anti-Fc<sub>Y</sub>RI murine mAb (M22 and M32.2) and a human IgG1 mAb were performed. Unconjugated H22 and M22 competed equivalently for either the binding of fluoresceinated M22 or fluoresceinated H22

10 10 in the presence of excess human IgG which saturated the Fc binding sites on Fc<sub>Y</sub>RI. As expected, the anti-Fc<sub>Y</sub>RI antibody M32.2 which binds to a different site on Fc<sub>Y</sub>RI than M22 (Guyre, P.M. et al., *J. Immunol.* 143:1650) was also unable to compete with the M22-FITC. In addition, the inhibition of H22-FITC by H22 and not by an irrelevant human IgG1 mAb confirmed the specificity of Fc<sub>Y</sub>RI binding via the V regions of H22.

15 15 H22, but not M22, was able to compete for Fc mediated binding to Fc<sub>Y</sub>RI by a fluoresceinated human IgG1. This experiment demonstrated that the Fc portion of H22 but not M22 bound to the Fc binding domain of Fc<sub>Y</sub>RI. This is consistent with the ability of the Fc portion of human IgG1 antibodies, but not murine IgG1, to bind Fc<sub>Y</sub>RI with high affinity.

20 20 Since the humanization of M22 was primarily to increase its immunotherapeutic potential, the binding activity of H22 to monocytes and cytokine-activated neutrophils was determined in the presence of human serum. H22-FITC bound with similar affinity to Fc<sub>Y</sub>RI on monocytes in the presence or absence of human serum. In 25 contrast, the Fc-mediated binding of an irrelevant human IgG-FITC was completely inhibited by human serum. Likewise, H22-FITC bound with similar affinity to IFN- $\gamma$ -treated neutrophils in the absence and in the presence of human serum. Collectively, the data demonstrated that H22 binds both via its V regions to a site distinct from the Fc binding domain and via its Fc region to the ligand binding domain of Fc<sub>Y</sub>RI. The former binding 30 30 activity effectively overcomes antibody blockade of human IgG1.

Functional Activity of H22 BsAb

The foremost application of anti-Fc<sub>Y</sub>RI antibodies for immunotherapy is the development of BsAbs which link Fc<sub>Y</sub>RI-bearing effector cells to a tumor cell, a virus, or a 35 35 virally-infected cell. Such BsAb have been developed with M22; therefore, a comparison was made of the ability of the M22 anti-tumor BsAb (520C9xM22) and a corresponding H22

BsAb (520C9xH22) to mediate cytotoxicity. These BsAbs consisted of H22 or M22 Fab' chemically conjugated to the Fab' of an anti-HER2/*neu* antibody (520C9), and thus were specific for the effector cell trigger molecule Fc<sub>Y</sub>RI and the tumor antigen.

5 Comparison of M22-derived and H22-derived BsAbs was done by ADCC assays. M22- and H22-derived BsAbs mediated the killing of HER2/*neu* overexpressing SKBR-3 cells. Both the murine and humanized BsAbs exhibited similar levels of lysis of antigen bearing target cells. In addition, both BsAb retained ADCC activity in the presence of human serum, while excess M22 F(ab')<sub>2</sub> resulted in complete inhibition of killing. Taken  
10 together these results show that the H22 BsAb-induced lysis is mediated through the M22 epitope and that the ADCC is Fc<sub>Y</sub>RI specific.

Finally, the ability of H22 and M22 to stimulate superoxide production by the monocyte-like cell line U937 was evaluated. M22, which binds to the Fc<sub>Y</sub>RI only by its V  
15 regions, induced a very low level oxygen burst, presumably because it is unable to cross-link the receptor efficiently. However, H22, which can cross-link Fc<sub>Y</sub>RI by binding as a ligand via its Fc domain and, additionally, as an antibody via its Fv, induced a more substantial release of superoxide.

20 **Example 2: Generation of a Functional H22-Epidermal Growth Factor Fusion Protein**

**Materials and Methods**

**Expression vectors and cloning**

25 Expression vectors for the genomic clones of the heavy (pSVgpt) and light (pSVhyg) chains of H22 are as described in International Patent Application Publication Number: WO 94/10332 entitled, *Humanized Antibodies to Fc Receptors for Immunoglobulin G on Human Mononuclear Phagocytes*. For the Fab-ligand fusion construct, it was unnecessary to alter the light chain. For the heavy chain, however, the CH2 and CH3  
30 domains had to be removed and replaced with the coding sequences of the ligands. The heavy chain vector contains two *Bam*HI sites, one in the intron between VH and CH1, and the other just downstream of CH3. Using the *Bam*HI restriction sites, DNA encoding the constant domains were replaced by a truncated version encoding only CH1 and most of the hinge. To do this, the polymerase chain reaction (PCR) was utilized to engineer the new C-  
35 terminus of the heavy chain fragment with the alterations shown in Figure 1.

The construct shown in Figure 1 [C], consisting of a translation termination codon downstream of the cloning restriction sites, *Xba*I and *Nol*I, and upstream of a *Bam*HI site which was used to clone the new PCR generated CH1 fragment downstream of VH, was

used to generate the fusion protein constructs. The cloning sites, which are located downstream of most of the hinge in order to retain flexibility between the Fd and ligand domains, was used to insert DNA encoding EGF or other ligands. Also, the single Cys residue has been retained from the previous construct to allow conjugation for the formation 5 of dimeric molecules.

DNA encoding the ligands were amplified by PCR to have a *Xba*I site on the N-terminus and a *Not*I site on the C-terminus of the coding region, and then inserted in the proper reading frame into the same sites of the newly engineered H22 heavy chain truncated 10 fragment described above. cDNA encoding epidermal growth factor (EGF) was obtained from the ATCC (#59957). Only DNA encoding the 53 amino acid residues of mature EGF out of the approximately 1200 residue precursor was cloned beginning with Asn 971 and ending with Arg 1023 (Bell, G.I., Fong, N.M., Stempien, M.M., Wormsted, M.A., Caput, D., Ku, L., Urdea, M.S., Rall, L.B. & Sanchez-Pescador, R. Human Epidermal Growth Factor 15 Precursor: cDNA Sequence, Expression In Vitro and Gene Organization. *Nucl. Acids Res.* 14: 8427-8446, 1986.).

#### Expression

The murine myeloma NSO (ECACC 85110503) is a non-Ig synthesizing line 20 and was used for expression of the fusion proteins. The final expression vector, a pSVgpt construct with DNA encoding H22 Fd fused in frame to EGF (shown in Figure 2) was transfected by electroporation using a BioRad Gene Pulser to NS0 which had been previously transfected with the pSVhyg construct containing DNA encoding H22 light chain. These polypeptides were expressed by an Ig promoter and Ig enhancer present in the vectors, and 25 secreted by the mAb 22 heavy chain signal peptide located on the N-terminus of the constructs. One or two days after transfection, mycophenolic acid and xanthine were added to the media to select for cells that took up the DNA. Individual growing colonies were isolated and subcloned after binding activity was demonstrated by ELISA.

#### Purification

Cells expressing the H22-EGF fusion protein were subcloned and expanded. The fusion protein-expressing clone was expanded and grown in spinner cultures and the supernatant was clarified and concentrated. Small scale purification was performed by affinity chromatography on an anti-human kappa chain affinity column (Sterogene, Carlsbad, 35 CA). The purified protein was analyzed by SDS-PAGE on a 5-15% acrylamide gradient gel under nonreducing conditions. Figure 3 is a schematic representation of the generation of anti-Fc receptor-ligand fusion proteins.

Bispecific flow cytometry

To show that the fusion protein is capable of binding both Fc<sub>Y</sub>RI and EGFR simultaneously, a flow cytometric assay has been developed (Figure 4). In this assay 5 different concentrations of H22-EGF fusion protein or the bispecific antibody, BsAb H447 (H22 X H425, a humanized version of the murine monoclonal antibody M425, which binds EGFR at the ligand binding site (E. Merck) was incubated with A431 cells, a cell line which expresses the EGF receptor (EGFR) (ATCC, Rockville, MD). After washing, a supernatant containing a fusion protein consisting of the extracellular domain of Fc<sub>Y</sub>RI and the Fc portion 10 of human IgM was added. Finally, a Phycoerythrin (PE)-labeled mAb (32.2), that binds Fc<sub>Y</sub> RI at a site that is distinct from that bound by mAb 22, was added. The cells were then analyzed by FACSCAN. Alternatively, binding to EGFR was blocked by excess (100 µg/ml) 15 whole murine mAb 425 (E. Merck), and binding of bsAb or fusion protein was detected by PE-labeled anti-human IgG.

15

ADCC

ADCC mediated by the fusion protein was determined using a <sup>51</sup>Cr killing assay. The EGFR overexpressing cell line, A431, was used as targets for lysis by human 20 monocytes cultured in  $\gamma$ -interferon (IFN- $\gamma$ ) for 24 hours. Targets were labeled with 100 µCi of <sup>51</sup>Cr for 1 hour prior to combining with effector cells and antibodies in a U-bottom microtiter plate. After incubation for 5 hours at 37°C supernatants were collected and 25 analyzed for radioactivity. Cytotoxicity was calculated by the formula: % lysis = (experimental CPM - target leak CPM/detergent lysis CPM - target leak CPM) X 100%. Specific lysis = % lysis with antibody - % lysis without antibody. The ability of the fusion 30 protein to mediate ADCC was compared with that of the respective BsAb. The assay was also performed in the presence of 25% human serum to demonstrate that IgG or other factors found in human serum will not inhibit fusion protein-mediated ADCC.

Results

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Purification

NS0 cells expressing the H22 kappa chain were transfected with the H22-EGF heavy chain construct and clones selected for resistance to mycophenolic acid and xanthine were expanded and the fusion protein was affinity-purified from the supernatant on an anti-35 human kappa column (Sterogene, Carlsbad, CA). The purified protein was analysed by SDS-PAGE. The purified protein migrated at an apparent molecular weight of 50-55 kDa, indicating that the fusion protein is expressed as a monomer, not a disulfide-linked dimer. In addition, a band was seen at an apparent molecular weight of 25kDa and is probably free light chain.

Binding specificity

To demonstrate that the fusion protein could bind Fc $\gamma$ RI and EGFR simultaneously a bispecific FACS assay was devised. Figure 5 shows that both the 5 chemically-linked, fully-humanized BsAb H447 (H22 (anti-Fc $\gamma$ RI) x H425), which was made as described in the following Example 3, and the H22-EGF fusion protein bound EGFR on A431 cells and soluble Fc $\gamma$ RI simultaneously in a dose-dependent fashion.

The EGFR-specificity of the fusion protein was demonstrated by the ability of 10 the murine mAb, M425, which binds EGFR at the ligand binding site, to inhibit fusion protein or H22 x H425 binding. Various concentrations of either the BsAb H447, or of the H22-EGF fusion protein were incubated with A431 cells in either the presence or absence of an excess of M425. Figure 6 shows that binding of both the BsAb and the fusion protein were inhibited by M425, demonstrating the specificity of the fusion protein for EGFR.

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ADCC

The ability of the fusion protein to mediate ADCC was analyzed using A431 cells as targets. Human monocytes cultured for 24 hours in the presence of IFN- $\gamma$  were used as effector cells. Figure 7 demonstrates the whole antibody, H425, the BsAb H447 (H22 x 20 H425) and the fusion protein mediated dose-dependent lysis of A431 cells. Figure 8 demonstrates that while ADCC mediated by the whole antibody is inhibited by 25% human serum (25%HS), ADCC mediated by the fusion protein was not inhibited by human serum and, in this particular experiment, fusion protein-mediated ADCC was enhanced by human serum. These results support the clinical utility of these molecules by demonstrating that the 25 fusion protein was capable of killing EGFR-overexpressing cells, even in the presence of Fc $\gamma$  RI-expressing effector cells as would be present *in vivo*.

Growth inhibitory properties of H22-EGF fusion proteins

Although EGF acts to stimulate growth of normal cells that express receptors 30 for it, EGF also can act to inhibit growth of tumor cells that over-express EGF-R (Barnes, D.W. (1982) *J. Cell Biol.* 93:1, MacLeod, C.L. et al. (1986) *J. Cell. Physiol.* 127:175). The ability of EGF and the H22-EGF fusion protein to inhibit the growth of A431 cells was examined as follows.

2 x 10<sup>4</sup> A431 cells were added to six well plates in complete media alone or in 35 media containing various concentration of either EGF, H22-EGF, the Fab fragment of H22, or the F(ab')<sub>2</sub> fragment of H425. Viable cells were counted after seven days using a hemocytometer. The analyses were performed in duplicate and reported as means +/- standard deviations.

The results are presented in Figure 9. These results indicate that EGF and H22-EGF significantly inhibited cell growth in a dose dependent fashion. On the contrary, the F(ab')2 fragment of H425 which had some inhibitory activity only at high concentrations and the Fab fragment of H22 had no growth inhibiting activity.

5

Thus, H22-EGF is able to bind to both Fc<sub>Y</sub>RI and EGF simultaneously, indicating that the molecule had folded properly and had maintained the flexibility required to bind both receptors at the same time. Furthermore, H22-EGF inhibited proliferation of the EGF-R expressing tumor cell line, A431, indicating that, similar to EGF, the H22-EGF fusion protein is capable of signaling through the EGF-R. H22-EGF also mediates potent killing of A431 cells in the presence of Fc<sub>Y</sub>RI expressing effector cells. Thus, H22-EGF mediates both cytotoxic and cytostatic effects on EGF-R expressing cells. Administration of H22-EGF to a subject having a tumor will result in recruitment of the body's natural cytotoxic effector cells to mediate killing of the tumor cells by potentially three different modes - cytotoxicity, growth inhibition, and phagocytosis. Furthermore, in addition to cell mediated cytotoxicity of the tumor cells, the effector cells recruited by H22-EGF may also further augment anti-tumor immunity by secreting inflammatory cytokines and/or by processing and presenting tumor antigens to tumor specific T cells.

20 **Example 3: H22-Heregulin (H22-gp30) Fusion Protein Mediates Tumor Cell Killing**

Heregulin (HRG) is a ligand for the HER3 and HER4 molecules. Both of these receptors may form heterodimers with HER2, a molecule which is overexpressed in some breast cancer cells. The affinity of HRG for HER3 and HER4 increases significantly when these molecules form heterodimers with HER2. This example demonstrates that a bispecific molecule, comprising heregulin and a binding specificity for the Fc<sub>Y</sub>RI inhibits growth of a tumor cell line and mediates fusion protein dependent cytotoxicity of these cells in the presence of Fc<sub>Y</sub>RI-bearing cytotoxic effector cells.

The H22-heregulin fusion protein was constructed in the same manner as the H22-EGF fusion protein described in Example 2. Briefly, genomic DNA encoding the Fd fragment of humanized anti-Fc<sub>Y</sub>RI mAb, H22, was fused to cDNA encoding the EGF domain of the  $\beta$ 2 form of HRG. The amino acid sequence of the H22-HRG fusion protein (SEQ ID NO: 4) is shown in Figure 10. This fusion protein comprises amino acids 171-239 of the heregulin  $\beta$ 2 shown in U.S. Patent Number 5,367,060. Other portions of heregulin  $\beta$ 2, as well as portions of other heregulin molecules, such as those disclosed in U.S. Patent Number 5,367,060 can also be used. The resulting H22Fd-HRG expressing vector was transfected into a myeloma cell line previously transfected with a vector containing DNA encoding the H22 kappa light chain. The resultant fusion protein was expressed predominantly as a monomer, even though the protein contains a free Cys residue in the hinge region of the H22

Fab component. Flow cytometry showed that this fusion protein was able to bind to the HER2 overexpressing tumor cell line, SKBR-3, as well as to Fc<sub>Y</sub>R-expressing cells.

5 To test the biological activity of the H22Fd-HRG fusion protein, supernatant from the myeloma cells expressing this fusion protein was diluted three fold or thirty fold and added to PC-3 cells or SKBR-3 tumor cells expressing HER2, HER3, and HER4 in the presence of IFN-treated monocytes at a ratio of 100:1 monocytes to target tumor cells. The monocytes were treated with IFN- $\gamma$  and the target cells were labeled with <sup>51</sup>Cr as described in Example 2. The % of specific lysis was calculated as indicated in Example 2. The results 10 are presented in Figure 11. The results indicate that about 45% of SKBR3 cells and up to about 49% of PC-3 cells are lysed upon incubation of the cells with the supernatant diluted 3 fold.

15 This fusion protein inhibits growth of SKBR-3 tumor cells and mediates fusion protein dependent cytotoxicity of these cells in the presence of Fc<sub>Y</sub>RI-bearing cytotoxic effector cells. Thus, the results of this example show that an anti-Fc<sub>Y</sub>RI-heregulin fusion protein can mediate anti-tumor cytotoxic activities under physiologic conditions and indicate that such a fusion protein will have therapeutic utility in the treatment of various cancers.

20 **Example 4: H22-Bombesin Fusion Protein Mediates Tumor Cell Killing**

25 The H22-bombesin fusion protein was constructed similarly to the H22-EGF fusion protein described above. However, since bombesin is a short peptide (14 amino acid residues), instead of amplifying cDNA encoding bombesin using PCR technology, DNA oligomers encoding the sense and anti-sense strands of bombesin were hybridized to create the coding region. The amino acid sequence of the bombesin peptide fused to the carboxyl end of the heavy chain of the H22 antibody is the following:

-Gln-Arg-Leu-Gly-Asn-Gln-Trp-Ala-Val-Gly-His-Leu-Met-Gly (SEQ ID NO: 5)

30 which corresponds to amino acids 2-14 of bombesin (Anastasi et al. (1971) *Experientia* 27:166) and contains an additional glycine residue at the carboxyl end of the peptide. The oligomers had overlapping ends that did not hybridize but instead created sticky ends for a *Xhol* site on the N-terminus and a *NotI* site on the C-terminus such that it could be cloned into the H22 heavy chain expression vector described above.

35 The biological activity of the H22-bombesin fusion protein on tumor cell killing was investigated as described above for the H22-EGF and H22-heregulin fusion proteins. Briefly, PC-3 tumor cells bearing bombesin receptors were labeled with <sup>51</sup>Cr and incubated with monocytes and various concentrations of H22-fusion protein, and fusion

protein dependent lysis was determined as described above. The results, shown in Figure 12, indicate that the target cells are lysed and that the level of target cell lysis increases proportionally with the amount of fusion protein added to the assay.

5                   Fusion proteins having H22 as one binding entity and CD4 (AIDS Repository) or gpl20 (AIDS Repository) as a second binding entity were also produced.

**Example 5: Production of Bispecific Antibodies From Modified Humanized Antibody Fragments**

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**Materials and Methods**

**Expression vectors and cloning**

15                   Expression vectors for the genomic clones of the heavy (pSVgpt) and light (pSVhyg) chains of H22 were as described in International Patent Application Publication Number: WO 94/10332 entitled, *Humanized Antibodies to Fc Receptors for Immunoglobulin G on Human Mononuclear Phagocytes*. For the Fab' construct, it was unnecessary to alter the light chain. For the heavy chain, however, the CH2 and CH3 domains had to be removed and replaced with a termination codon. The heavy chain vector contains two *Bam*HI sites, 20 one in the intron between VH and CH1, and the other just downstream of CH3. Using the *Bam*HI restriction sites, DNA encoding the constant domains were replaced by a truncated version encoding only CH1 and most of the hinge. To do this, The polymerase chain reaction (PCR) was utilized to engineer the new C-terminus of the heavy chain fragment with the alterations shown in Figure 1. Figure 1 [B] shows the alterations for generation of a 25 truncated single-sulphydryl version.

**Expression**

30                   The murine myeloma NSO (ECACC 85110503) is a non-Ig synthesizing line and was used for expression of the modified H22 antibody. The final expression vector, a pSVgpt construct with DNA encoding H22 Fd was cotransfected with the pSVhyg construct containing DNA encoding H22 light chain by electroporation using a BioRad Gene Pulser. These polypeptides were expressed by an Ig promoter and Ig enhancer present in the vectors, and secreted by the mAb 22 heavy chain signal peptide located on the N-terminus of the constructs. One or two days after transfection, mycophenolic acid and xanthine were added 35 to the media to select for cells that took up the DNA. Individual growing colonies were isolated and subcloned after Fc $\gamma$ RI binding activity was demonstrated.

Purification

5 The single sulphydryl form of the H22 antibody and the whole H425 (anti-EGFR) antibody were produced by *in vitro* cultivation of the respective transfected NSO cells. The H425 was purified by protein A affinity chromatography. The single sulphydryl form of the antibody H22 was purified by ion exchange chromatography using Q-Sepharose followed by SP-Sepharose (Pharmacia, Piscataway, NJ). The purity of the single sulphydryl form of the H22 antibody was assessed by SDS-PAGE.

Generation of bispecific antibody (BsAb)

10 BsAb was constructed using the method of Glennie et al. (Glennie, M.J. et al., 1987), Preparation and performance of bispecific  $F(ab')_2$ , antibody containing thioether-linked Fab' gamma fragments, *J. Immunol.*, 139:2367). The  $F(ab')_2$  of H425 was generated by limited pepsin proteolysis in 0.1M citrate buffer, pH 3.5 and the  $F(ab')_2$  purified by ion exchange chromatography. The mAbs were reduced by addition of 20 mM 15 mercaptoethanolamine (MEA) for 30 minutes at 30°C. The Fab' fragments were applied to a Sephadex G-25 column equilibrated in 50mM sodium acetate, 0.5mM EDTA, pH 5.3 (4°C). Ortho-phenylenedimaleimide (o-PDM, 12mM) dissolved in dimethyl formamide and chilled in a methanol/ice bath was added (one half volume) to the H22 Fab' and incubated for 30 minutes on ice. The Fab'-maleimide was then separated from free o-PDM on Sephadex G-25 20 equilibrated in 50mM Na Acetate, 0.5mM EDTA, pH 5.3 (4°C). For preparation of the BsAbs, the H22 Fab'-maleimide was added to the H425 Fab' at a 1.2:1 molar ratio. The reactants were concentrated under nitrogen to the starting volume using a Diaflo membrane in an Amicon chamber (all at 4°C). After 18 hours the pH was adjusted to 8.0 with 1M Tris-HCl, pH 8.0. The mixture was then reduced with 10mM MEA (30 minutes, 30°C) and 25 alkylated with 25 mM iodoacetamide. The bispecific  $F(ab')_2$  was separated from unreacted Fab's and other products by a Superdex 200 (Pharmacia, Piscataway, NJ) column equilibrated in PBS.

Bispecific flow cytometry

30 To show that BsAb generated by the o-PDM method as well as that generated by the DTNB method are capable of binding both Fc $\gamma$ RI and EGFR simultaneously, a flow cytometric assay has been developed (Figure 13). In this assay different concentrations of the two BsAbs were incubated with A431 cells, a cell line which expresses the EGF receptor (EGFR). After washing, a supernatant containing a fusion protein consisting of the 35 extracellular domain of Fc $\gamma$ RI and the Fc portion of human IgM was incubated with the cells. Finally, the cells were incubated with a FITC-labeled anti-human IgM-specific antibody. The cells were then analyzed by FACSCAN.

ADCC

BsAb-mediated ADCC was determined using a  $^{51}\text{Cr}$  killing assay. The EGFR overexpressing cell line, A431, was used as targets for lysis by human monocytes cultured in  $\gamma$ -interferon for 24 hours. Targets were labeled with 100  $\mu\text{Ci}$  of  $^{51}\text{Cr}$  for 1 hour prior to combining with effector cells and antibody in a flat-bottomed microtier plate. After incubation for 16 hours at 37°C supernatants were collected and analyzed for radioactivity. Cytotoxicity was calculated by the formula: % lysis = (experimental CPM - target leak CPM/detergent lysis CPM - target leak CPM) X 100%. Ab-dependent lysis = % lysis with antibody - % lysis without antibody.

10

ResultsPurification

NSO cells were cotransfected with the truncated H22 heavy chain construct and the intact kappa chain construct. Clones selected for resistance to mycophenolic acid and xanthine were expanded and the protein was purified from the supernatant by Q-Sepharose followed by SP-Sepharose ion exchange chromatography. The purified protein was analyzed by SDS-PAGE. The purified protein migrated at an apparent molecular weight of 50kDa, indicating that the protein is expressed as a monomer, not a disulfide-linked dimer.

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Construction and characterization of a BsAb composed of single sulphydryl H22 linked to Fab' of H425 (anti-EGFR)

A BsAb was constructed where the single sulphydryl form of H22 was linked to the Fab' fragment of H425, a humanized anti-EGFR mAb. The BsAb was generated using o-PDM as a linker by the method of Glennie et al. (Glennie, M.J. et al., (1987), Preparation and performance of bispecific F(ab' gamma)<sup>2</sup>, antibody containing thioether-linked Fab' gamma fragments, *J. Immunol.*, 139:2367). The activity of this BsAb was compared to one generated by the DTNB method using Fab' fragments made from pepsin digestion and reduction of whole H22. To demonstrate that these BsAbs could bind Fc $\gamma$ RI and EGFR simultaneously a bispecific FACS assay was devised. Figure 14 shows that both the o-PDM-linked BsAb and the BsAb made by the DTNB method bound EGFR on A431 cells and soluble Fc $\gamma$ RI simultaneously in a dose-dependent fashion.

The ability of the two BsAbs to mediate ADCC was analyzed using A431 cells as targets. Human monocytes cultured for 24 hours in the presence of IFN- $\gamma$  were used as effector cells. Figure 15 demonstrates the two BsAbs mediated dose-dependent lysis of A431 cells in a comparable fashion. These results demonstrated that BsAb generated from the truncated, single sulphydryl form of H22 was capable of killing EGFR-overexpressing cells in the presence of Fc $\gamma$ RI-expressing effector cells.

Example 6: Production of Trivalent Antibodies

Materials and Methods

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Cell lines and antibodies. M22, 520C9, H425, SKBR3 and A431

M22 and 520C9 were purified from hybridoma supernatant by ion exchange chromatography (Pharmacia, Piscataway, NJ) and 520C9 was further purified by protein A affinity chromatography (Pharmacia, Piscataway, NJ). H425 was purified from hybridoma supernatant by protein A affinity chromatography (Pharmacia, Piscataway, NJ). The M22- and 520C9-producing murine hybridoma were described previously (Guyre et al., (1989) Monoclonal antibodies that bind to distinct epitopes on Fc<sub>γ</sub>RI are able to trigger receptor function, *J. Immunol.* **143**:5, 1650-1655; Frankel et al., (1985) Tissue distribution of breast cancer-associated antigens defined by monoclonal antibodies, *J. Biol. Response Modifiers*, **4**:273-286). The murine myeloma NSO (ECACC 85110503) is a non-Ig synthesizing line and was used for the expression of the humanized mAb, H425 (Kettleborough et al., (1991) Humanization of a mouse monoclonal antibody by CDR-grafting: the importance of framework residues on loop conformation, *Protein Eng.*, **4**:773). SKBR-3, (ATCC, Rockville, MD) a human breast carcinoma cell line that overexpresses the HER2/neu protooncogene, and A431 (ATCC, Rockville, MD), a human squamous carcinoma cell line that overexpresses EGFR (ATCC, Rockville, MD) were cultivated in Iscove's Modified Dulbecco's Medium (IMDM, Gibco, Grand Island, NY).

Neutrophil preparation

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Neutrophils are separated from mononuclear cells by ficoll hypaque (Pharmacia, Piscataway, NJ) gradient separation. To up-regulate F<sub>cγ</sub>RI, neutrophils are treated with cytokines. Neutrophils are cultured for 24-48hrs (37°C, 5% CO<sub>2</sub>) in AIM V media (Gibco, Grand Island, NY) containing 2.5% normal human serum type AB (Sigma, St. Louis, MO), 50 ng/ml G-CSF (Kindly provided by R. Repp, U. of Erlanger, Germany) and 30 100 IRU/ml IFN-γ.

Conjugation method

35

BsAb were constructed using the method of Glennie et al (Glennie, M.J. et al., (1987), Preparation and performance of bispecific F(ab')<sup>2</sup>, antibody containing thioether-linked Fab' gamma fragments, *J. Immunol.*, **139**:2367). mAbs M22, 520C9 (anti-HER2/neu, 33), and H425 (anti-EGFR) antibodies were produced by *in vitro* cultivation of the respective hybridoma cells. The F(ab')<sup>2</sup> of each antibody were generated by limited pepsin proteolysis in 0.1M citrate buffer, pH 3.5 and the F(ab')<sup>2</sup> purified by ion exchange chromatography. mAbs M22 and H425 were reduced to Fab' by addition of 20 mM

mercaptoethanolamine (MEA) for 30 minutes at 30°C. The Fab' fragments were applied to a Sephadex G-25 column equilibrated in 50mM Na Acetate, 0.5mM EDTA, pH 5.3 (4°C). Ortho-phenylenedimaleimide (o-PDM, 12mM) dissolved in dimethyl formamide and chilled in a methanol/ice bath was added (one half volume) to the murine 22 Fab' and incubated for 5 30 minutes on ice. The Fab'-maleimide was then separated from free o-PDM on Sephadex G-25 equilibrated in 50mM Na Acetate, 0.5mM EDTA, pH 5.3 (4°C). For preparation of the BsAbs, the M22 Fab'-maleimide was added to the H425 Fab' at a 1:1 molar ratio. The reactants were concentrated under nitrogen to the starting volume using a Diaflo membrane in an Amicon chamber (all at 4°C). After 18 hours the pH was adjusted to 8.0 with 1M Tris-10 HCl, pH 8.0. The mixture was then reduced with 10mM MEA (30 minutes, 30°C) and alkylated with 25 mM iodoacetamide. The bispecific F(ab')<sub>2</sub> as separated from unreacted Fab's and other products by a Superdex 200 (Pharmacia, Piscataway, NJ) column equilibrated in phosphate buffered saline (PBS). The BsAb M22 x 520C9 was made in a similar fashion except that 520C9 was used instead of H425.

15

Trispecific antibody composed of M22 x H425 x 520C9 was made in two stages (Figure 16). In the first stage, M22 was linked to H425 as described above to create the M22 x H425 BsAb except that rather than a final reduction and alkylation, the reactants were treated with DTNB to block the remaining free sulfhydryl groups. The bivalent BsAb 20 was purified by gel filtration on a Superdex 200 column, reduced to F(ab')<sub>2</sub>(SH) and mixed in a 1:1 molar ratio with o-PDM-treated 520C9. The resulting trispecific F(ab)3 was purified on a Superdex 200 column. The TsAb was analyzed by HPLC size exclusion chromatography using a TSK 3000 column (ToJo Haas, Japan). Using the same procedure as above another TsAb comprising m22 Fab' x 32.2 Fab' x m22 Fab' has been constructed.

25

#### Bispecific flow cytometry

The TsAb can bind to EGFR and F<sub>cyt</sub>RI simultaneously or to HER2/neu and F<sub>cyt</sub>RI simultaneously. Either A431 cells (high EGFR-expressing cells) or SKBR-3 cells (high HER2/neu-expressing cells) were incubated with various concentrations of BsAbs 30 (M22 x 520C9 or M22 x H425) or with the TsAb, M22 x H425 x 520C9. The cells were washed and then incubated with the soluble F<sub>cyt</sub>RI. Soluble F<sub>cyt</sub>RI binding was detected with mAb 32.2-FITC which binds F<sub>cyt</sub>RI at a site that is distinct from the 22 binding site. The cells were then analyzed by FACSCAN.

35

#### ADCC

Either SKBR-3 cells or A431 cells were used as targets for lysis by cytokine activated neutrophils. Targets were labeled with 100 $\mu$ Ci of <sup>51</sup>Cr for 1 hour prior to combining with neutrophils and antibodies in a U-bottom microtiter plate. After incubation for 16 hours at 37°C supernatants were collected and analyzed for radioactivity. Cytotoxicity

was calculated by the formula: % lysis = (experimental CPM - target leak CPM/detergent lysis CPM - target leak CPM) X 100%. Specific lysis = % lysis with antibody - % lysis without antibody. Assays were performed in triplicate.

5

#### Fc<sub>Y</sub>RI Modulation Assay

The M22 x 32.2 x M22 BsAb was used for modulation of Fc<sub>Y</sub>RI on monocytes in whole blood. The assay procedure is shown in the enclosed flow chart (see Figure 23A). Figure 23B shows that treatment with 10 $\mu$ g/mL of this BsAb decreased the Fc<sub>Y</sub>RI expression on monocytes to approximately 50% of the level prior to BsAb treatment.

10

#### Results

##### Construction and biochemical characterization of the TsAb

TsAb was made according to the flow chart depicted in Figure 16. In the first 15 stage of the procedure, M22 was coupled to H425, treated with DTNB, and the resulting bispecific F(ab')<sub>2</sub> purified by gel filtration. In the second stage, this bispecific F(ab')<sub>2</sub> was reduced and mixed with o-PDM-treated 520C9 Fab' resulting in the TsAb, M22 x H425 x 520C9. This TsAb is depicted schematically in Figure 17. In this figure, Fab'-A represents M22, Fab'-B represents H425, and Fab'-C represents 520C9.

20

##### Binding (Bs FACS)

To demonstrate that the TsAb, M22 x H425 x 520C9, could bind Fc<sub>Y</sub>RI and HER2/neu simultaneously a bispecific FACS assay was devised. This assay is depicted schematically in Figure 18A. Figure 19 shows that both the TsAb bound HER2/neu on 25 SKBR-3 cells and soluble Fc<sub>Y</sub>RI simultaneously in a dose-dependent fashion. The BsAb, M22 x H425, generated negligible signal in this assay over a wide range of concentrations. To demonstrate that the TsAb, M22 x H425 x 520C9, could bind Fc<sub>Y</sub>RI and EGFR simultaneously a similar assay was devised using the EGFR-overexpressing cell line, A431, in the case. This assay is depicted schematically in Figure 18B. Figure 20 shows that both 30 the TsAb and the BsAb, M22 x H425, bound EGFR on A431 cells and soluble Fc<sub>Y</sub>RI simultaneously in a dose-dependent fashion. The BsAb, M22 x 520C9, generated negligible signal in this assay over a wide range of concentrations.

##### ADCC

35 The ability of the TsAb to mediate ADCC was analyzed using either SKBR-3 or A431 cells as targets. Human neutrophils cultured for 24-48 hours in the presence of IFN- $\gamma$  and G-SF were used as effector cells. Figure 21 demonstrates the both the BsAb, M22 x 520C9, and the TsAb, M22 x H425 x 520C9, mediated lysis of SKBR-3 cells, whereas the

BsAb, M22 x H425, did not. On the other hand, Figure 22 demonstrates the BsAb, M22 x H425, and the TsAb, mediated lysis of SKBR-3 cells, whereas the BsAb, M22 x 520C9, did not. These results demonstrated that the TsAb was capable of killing both HER2/neu and EGFR-overexpressing cells in the presence of F<sub>cγ</sub>RI-expressing effector cells.

5

The trispecific antibody described above included M22, the murine version of the anti-Fc<sub>γ</sub>RI mAb. Such a trispecific antibody could be constructed using the single-sulphydryl form of the humanized anti-Fc<sub>γ</sub>RI mAb, H22. The only difference being that single-sulphydryl form is secreted as a F(ab')<sub>2</sub> fragment of this antibody. The single-sulphydryl form is purified from culture supernatants utilizing ion exchange chromatography using Q-Sepharose followed by SP-Sepharose (Pharmacia, Piscataway, NJ). Once the single-sulphydryl form of H22 is purified, the creation of a trispecific antibody using this reagent would be identical to that described above using the F(ab')<sub>2</sub> fragment of M22.

10

**15                   Example 7: Enhanced Antigen Presentation with H22-antigen Fusion Proteins**

20

This example demonstrates that (a) antigenic peptides genetically grafted onto the constant region of an anti-Fc<sub>γ</sub>RI antibody are significantly more efficient in antigen presentation of the antigen and T cell stimulation compared to the antigen alone, and (b) that antagonistic peptides genetically grafted onto the constant region of an anti-Fc<sub>γ</sub>RI are significantly more efficient in inhibiting T cell stimulation compared to the antagonistic peptide alone. Thus, such fusion proteins will effectively increase the delivery of peptides to antigen presenting cells (APCs) *in vivo* and will be useful in various therapeutic methods.

25

**Materials and Methods**

**Reagents**

30

AIM V (GIBCO, Grand Island, NY) was used as culture medium. Tetanus Toxoid (TT) was purchased from ACCURACTE CHEMICAL CO. (Westbury, NY). Sterile and low-endotoxin F(ab')<sub>2</sub> fragment of mouse anti-Fc<sub>γ</sub>RI mAb 22 and the bispecific Ab, MDXH210 (consisting of Fab' of humanized Ab 22 chemically linked to Fab' of anti-Her2/neu tumor Ag mAb 520C9) were provided by MEDAREX, INC. (Annandale, NJ). The universal Th epitope of TT, TT830-844 (QYIKANSKFIGITEL (SEQ ID NO: 6), termed as TT830 hereafter) (Valmori D. et al. (1994) *J. Immunol.* 152:2921-29) and the mutant form of this epitope, TT833S (QYISANSKFIGITEL (SEQ ID NO: 9), lysine at position 833 changed into serine) were synthesized and purified to >95% by PEPTIDOGENIC CO. (Livermore, CA). Another universal Th epitope of TT, TT947-967 (FNNFTVSF WLRVPKVSASHLE (SEQ ID NO: 12), referred to as TT947 hereafter), (>80% pure) (Valmori D. *supra*) was used

as a control peptide in this study. Commercially available human IgG for intravenous injection (IVI) was used in blocking experiments.

Cells

5 The monocytic cell line, U937, which expresses Fc<sub>Y</sub>RI, was obtained from the ATCC. The method of generating CD4<sup>+</sup>, peptide TT830-specific T cells was modified from a previously described protocol for TT-specific T cell lines (Gosselin E.J. (1992) *J. Immunol.* 149:3477-81). Briefly, mononuclear cells were isolated from peripheral blood using Ficoll Hypaque. 150 x 10<sup>6</sup> mononuclear cells were stimulated in 50 ml of AIM V medium with 10  
10 µM TT830. After three days' incubation at 37°C in a 5% CO<sub>2</sub> incubator, non-attached (mostly non-specific cells) were removed by washing the flask 1X with 10 ml of HEPES-buffered RPMI 1640; specific T cell colonies together with adherent monocytes remained in the flask. Fifty ml of AIM V plus 20 U/ml of human IL-2 (IMMUNEX, Seattle WA), and 1  
15 ml (2%, final concentration) pooled human serum were added back to the flask. After 10-14 days of total incubation time, T cells were harvested and dead cells were pelleted through Ficoll Hypaque, yielding a highly enriched population (95-98%) of viable CD4<sup>+</sup>, Ag-specific T cells. The T cells were confirmed to be specific for TT830 peptide as shown in Fig. 3.  
20 Large quantities of monocytes were purified from leukapheresis packs using the cold aggregation method (Mentzer S.J. et al. (1986) *Cell. Immunol.* 101:132) which resulted in 80-90% purity. Both monocytes and T cells were frozen in aliquots for future use and were shown to function normally after being thawed.

Ag presentation assay

25 In proliferation assays, T cells (5 x 10<sup>4</sup>), irradiated monocytes (3000 rad, 10<sup>5</sup>/well), and various concentrations of peptide TT830 fusion protein Fab22-TT830 were incubated together in a final volume of 200 µl/well in flat-bottom 96-well tissue culture plates for 2 days. 10µl (1 µCi/well) <sup>3</sup>H-thymidine was then added to each well. After incubating overnight, plates were harvested and counted in a liquid scintillation counter. T cell proliferation was expressed as the mean counts/min (CPM) of three replicates  $\pm$  SD.  
30 Background CPM (T cells and monocytes without Ag) was subtracted from all the data points. Experiments with APL were done according to similar protocols reported by Sette et al. (De Magistris (1992) *Cell* 68:625). Briefly, for inhibition assays, irradiated monocytes were treated with various concentrations of TT833S or Fab22-TT833S overnight. 20nM TT830 and T cells were then added. After a further 2 days incubation, T cell proliferation was measured as described above. In "pre-pulsing" experiments, irradiated monocytes were pulsed with 20 nM TT830 4 h prior to the addition of 10 µM TT833S or 0.1µM Fab22-TT833S. After overnight incubation, T cells were then added. After a further 2 days incubation, T cells were stimulated with irradiated monocytes and TT833S or Fab22-TT833S for 1 day, recovered after centrifugation over Ficoll Hypaque, and restimulated with

monocytes and various concentrations of TT830 for 2 days. T cell proliferation was then measured by the incorporation of  $^3\text{H}$ -thymidine and the average CPM of three replicates was plotted. In some cases, the percentage of inhibition was calculated by the formula: % inhibition =  $(\text{CPM}_{\text{no inhibitor}} - \text{CPM}_{\text{inhibitor}})/\text{CPM}_{\text{no inhibitor}} \times 100$ . All experiments were 5 repeated at least three times.

#### Staining and Flow Cytometry

Staining procedures were adapted from those previously described (Gosselin E.J. et al. (1990) *J. Immunol.* 144:1817-22). Briefly, to individual wells of a 96-well plate at 10 4°C, 30  $\mu\text{l}$  of RPMI+ 1mg/ml BSA containing one of the proteins Fab22-TT830, Fab22-TT833S, or the BsAb MDXH210 at varying concentrations. After 1 h incubation at 4°C, plates were centrifuged, the supernatants discarded, and the cells washed three times with PBS/BSA at 4°C. Cells were then incubated for 1 h with 40  $\mu\text{l}$  /well of FITC-labeled F(ab')<sub>2</sub> goat anti-human IgG (JACKSON IMMUNORESEARCH LABORATORIES, INC. West 15 Grove, PA) followed by three washes with PBS/BSA and resuspended in PBS/BSA containing 1% paraformaldehyde (KODAK, Rochester, NY). Cells were then examined by FACScan (BECTON DICKINSON & CO., Mountain View, CA), and mean fluorescence intensity (MFI) was measured.

20 Cytokine measurement

Supernatants were collected from the 96-well plates of Ag presentation assays after 2 days stimulation and frozen until used. The levels of IFN- $\gamma$  and IL-4 from these samples were measured by specific ELISA. Ab pairs for the IFN- $\gamma$  and IL-4-specific ELISA were purchased from PHARMINGEN (San Diego, CA). ELISA assays were performed 25 according to the protocol provided by the manufacturer.

#### Generation of H22-TT peptide fusion proteins

In order to generate fusion proteins Fab22-TT830 and Fab22-TT833S, synthetic oligonucleotides encoding each peptide were separately engineered into the hinge 30 region in the heavy chain of humanized anti-Fc $\gamma$ RI mAb 22 (H22) according to the method set forth below.

#### Expression and cloning vectors

mAb 22 has been humanized by grafting its CDR regions into a human IgG1 framework (see above and Graziano R. F. et al. (1995) *J. Immunol.* 155:4996-5002). The expression vector for the genomic clone of the heavy chain (pSVgpt) of H22 was modified to allow incorporation of the coding sequence for other molecules, in this case, the TT peptides. The BamHII fragment of this vector containing CH1, hinge, and newly engineered XbaI and 35 NsiI cloning sites (see Figure 2) was inserted into the BamHII site of pUC19 to generate the

vector pUC19/H22CH1(X+N). This vector was used to clone oligonucleotide sequences encoding TT peptides, as described below.

The oligonucleotide sequences encoding the tetanus toxin (TT) peptides were  
5 designed to have a XhoI site on the N-terminus and a NotI site on the C-terminus of the  
coding region (Figure 24A). These oligonucleotides were synthesized and purified by  
GENOSYS Biotechnologies (The Woodlands, TX). The synthetic oligonucleotides were  
then annealed and ligated into the cloning vector pUC19/H22CH1(X+N). Clones which had  
incorporated the coding sequences for TT peptides were screened by restriction mapping.  
10 The BamHI fragment containing CH1, hinge, and TT830 or TT833S was then cut out of  
pUC19 and inserted into the expression vector which already contained VH. The final  
expression construct of H22 heavy chain fused with TT peptides is shown in Figure 24B.

Expression of the H22-TT fusion proteins

15 The murine myeloma NSO (ECACC 85110503) is a non-Ig synthesizing line  
and was used for expression of the H22-TT fusion proteins. First, NSO cells were transfected  
with the pSVhyg vector containing the H22 light chain coding sequence. The H22 light  
chain expressing NSO cells were then transfected with the expression vector construct  
containing the H22 H-chain Fd sequence fused in frame to the TT coding sequences (Figure  
20 24B). A BioRad Gene Pulser electroporation apparatus was used to carry out the transfection  
employing 200 v and 960  $\mu$ Farad. One or two days after transfection, mycophenolic acid  
(0.8  $\mu$ g/ml; SIGMA) and xanthine (2.5  $\mu$ g/ml; SIGMA) were added to the media to select  
transfectants which had successfully taken up the expression vectors. Individual colonies  
25 were isolated based on the binding activity of the culture supernatants to Fc $\gamma$ RI on U937 cells  
as demonstrated by flow cytometry. The positive colonies were subcloned by limiting  
dilution.

Purification of the Fab22-TT fusion proteins

Clone pW5 expressing the Fab22-TT830 fusion protein and clone pM4  
30 expressing the Fab22-TT833S fusion protein were expanded in roller bottle cultures. The  
supernatants were clarified and concentrated. Small scale purification was performed by  
affinity chromatography on an anti-H22 affinity column. SDS-PAGE analysis of a 5-10%  
acrylamide gradient gel under non-reducing conditions showed that fusion proteins were  
>90% pure and had a molecular weight of 50 kDa as expected. Protein concentration was  
35 determined by absorbance at 280 nm using the extinction coefficient of IgG Fab' = 1.53.

ResultsH22Fd-TT fusion proteins bind to U937 cells

The ability of the H22 fusion proteins, Fab22-TT830 and Fab22-TT833S, to bind to Fc $\gamma$ RI was examined first. A previously described bispecific Ab, MDXH210, which contains the same Fc $\gamma$ RI-binding component (Fab' of humanized mAb 22) (Valone F. H. et al. (1995) *J. Clin. Oncol.* 13:2281-92), was used as a positive control. Binding of fusion proteins and MDXH210 to U937 cells, which constitutively express Fc $\gamma$ RI, was measured by staining with FITC-labeled goat Ab specific for human IgG and flow cytometry. As indicated in Figures 24A and 24B, fusion proteins Fab22-TT830 and Fab22-TT833S bound to U937 cells in a dose-dependent manner similar to MDXH210. The binding of fusion proteins was completely blocked by murine anti-human Fc $\gamma$ RI mAb 22 F(ab')<sub>2</sub>, demonstrating the specificity of fusion proteins for Fc $\gamma$ RI.

15                   H22Fd-TT fusion protein enhances presentation of TT peptide by 100-1000 fold

The fusion protein, Fab22-TT830 was used in Ag presentation assays to determine whether the Th epitope, TT830, when expressed in the constant region of H22, could be effectively presented by monocytes to autologous T cells. As shown in Figure 26, about 1,000-fold less Fab22-TT830 was required than TT830 peptide alone to achieve the same level of T cell proliferation. In addition, Figure 26 shows that the presentation of Fab22-TT830 was about 10,000-fold more efficient than the presentation of the intact TT, suggesting that the enhanced presentation of Fab22-TT830 did not merely result from higher molecular weight nor increased stability of Fab22-TT830 as opposed to TT830 peptide. Another antigenic TT epitope, TT947, failed to stimulate the T cells, confirming that the T cells were specific for TT830 peptide. These results thus provide clear evidence that Th epitopes expressed in the constant region of H22 can be effectively and specifically presented.

30                   Blockade of Fc $\gamma$ RI on monocytes abrogates the enhancement of Ag presentation by the H22Fd-TT fusion protein

To directly determine whether the enhancement of peptide presentation through the use of the fusion protein is Fc $\gamma$ RI-mediated, binding of Fab22-TT830 to Fc $\gamma$ RI on Ag-presenting monocytes was blocked by treating monocytes with mAb 22 F(ab')<sub>2</sub> for 1 h prior to the addition of Fab22-TT830 or TT830 peptides. Enhancement of peptide presentation by the fusion protein was abrogated by mAb 22 F(ab')<sub>2</sub>, whereas presentation of TT830 was unaffected (Figure 27). The fact that the binding of mAb 22 F(ab')<sub>2</sub> to Fc $\gamma$ RI did not lead to an enhancement of the presentation of free peptides implies that binding of mAb 22 to Fc $\gamma$ RI alone did not alter the functional state of monocytes in a way that enhanced Ag

presentation. Therefore, linkage of the peptide to anti-Fc $\gamma$ RI Ab 22 appears to be necessary for the observed enhancing effects on Ag presentation, suggesting that the enhanced presentation is probably a result of efficient Ag capture through Fc $\gamma$ RI.

5

Enhancement of peptide presentation by H22 is not affected by the presence of human IgG

Under physiological conditions, the ligand-binding domain of Fc $\gamma$ RI is saturated by IgG which blocks efficient targeting of AgAb to this receptor. A unique advantage for using derivatives of mAb 22 to trigger Fc $\gamma$ RI function is that Ab 22 binds to an epitope outside the ligand binding domain. Therefore, functions triggered by mAb 22, such as ADCC, phagocytosis and Ag presentation are not inhibited by physiological levels of IgG (Gosselin E. J., *supra*, Guyre P. M. (1989) *J. Immunol.* 143:1650-55). Similarly, the enhanced presentation of the TT830 peptide using the fusion protein Fab22-TT830 was not inhibited by IgG (Figure 28), suggesting that H22-based fusion proteins is also an effective way to target peptide Ags to Fc $\gamma$ RI *in vivo*.

20

IFN- $\gamma$  and IL-4 production is increased following H22Fd-TT fusion protein-enhanced Ag presentation

Upon activation, T cells not only undergo clonal expansion through proliferation, but also produce cytokines such as IFN- $\gamma$  and IL-4 to exert their effector function of B cell differentiation and monocyte activation (Paul W. E. and Seder, R. A. (1994) *Cell* 76:241-251). Therefore, the production of IFN- $\gamma$  and IL-4 following H22 fusion protein-enhanced Ag presentation was examined. As shown in Figures 28A and 28B, both IFN- $\gamma$  and IL-4 production levels were enhanced by Fab22-TT830, especially at suboptimal Ag concentrations. However, in these experiments, the enhancement for cytokine production (about 20-fold) was less than that for T cell proliferation (about 600-fold).

30

Thus, Th epitopes expressed in the constant region of anti-Fc $\gamma$ RI mAb H22 can be effectively processed and presented by human monocytes, leading to enhanced T cell activation and cytokine production.

Presentation of APL, TT833S and Fab22-TT833S fails to stimulate T cell proliferation

Peptides containing one or two amino acid changes from native T cell epitopes, termed Altered Peptide Ligands (APL) by Allen and coworkers, have been shown to be agonists, partial agonists, or antagonists for T cell activation (Sette et al. (1994) *Ann. Rev. Immunol.* 12:413 and Evavold et al. (1993) *Immunol. Today* 14:602). Recognition of APL by specific T cells through TCR in some cases triggered partial signal transduction and resulted in (i) inhibition of T cell stimulation by superantigen (Evavold et al. (1994) *Proc.*

Natl. Acad. Sci. U.S.A. 91:2300), T cell anergy (Sloan-Lancaster et al. (1993) Nature 363:156 and Sloan-Lancaster et al. (1994) J. Exp. Med. 185:1195), or (iii) modulation of Th1/Th2 differentiation (Nicholson et al. (1995) Immunity 3:397; Pfeiffer et al. (1995) J. Exp. Med. 181:1569; and Windhagen et al. (1995) Immunity 2:373). Partial agonists have 5 been shown to stimulate some T cell functions such as IL-4 production by T cells, but not others such as T cell proliferation (Evabold et al. (1991) Science 252:1308). Partial agonists also can induce anergy. Certain APL do not trigger any detectable signaling events upon interaction with TCR, but can function as TCR antagonists to inhibit T cell proliferation in response to wild-type peptide antigen and are thus called TCR antagonists (De Magistris et 10 al. (1992) Cell 68:625 and Ruppert et al. (1993) Proc. Natl. Acad. Sci. USA 90:2671.

This example demonstrates that the peptide TT833S, an antagonist peptide for T cell epitope TT830 of tetanus toxin and Fab22-TT833S fail to stimulate T cell proliferation. As shown in Figure 30, even at doses as high as 100  $\mu$ M for TT833S and 1  $\mu$ M for Fab22-TT833S, no significant proliferation of TT830-specific T cells was observed. This indicates 15 that changing lysine to serine at position 833 of the TT830 peptide eliminated T cell reactivity of this T cell epitope. Additionally, when peptides TT830 and TT833S were simultaneously presented to TT830-specific T cells, T cell proliferation in response to TT830 was inhibited by TT833S in a dose-dependent fashion, showing that TT833S can function as 20 an antagonist for TT830-specific T cells (Figure 31).

*Fab22-TT833S is at least 100 times more effective than TT833S in inhibiting T cell activation*

This example compares the relative efficiency of TT833S and Fab22-TT833S 25 in inhibiting T cell proliferation in response to TT830. As shown in Figure 32, Fab22-TT833S was about 100 times more effective than TT833S in inhibiting TT830-stimulated T cell proliferation. This suggests that the APL, TT833S, when expressed in the constant region of mAb H22, can be correctly and effectively presented by APC. The increased antagonistic efficacy of fusion protein Fab22-TT833S on T cell proliferation probably 30 reflects more efficient Ag capture mediated by Fc $\gamma$ RI as compared to free peptides.

*Inhibition of T cell activation is mediated by competition for T cell receptor binding rather than for MHC Class II binding.*

The antagonist effects of APL TT833S and fusion protein Fab22-TT833S 35 might be through competition at the level of MHC-binding or TCR-binding, or both. To gain insight into the mechanisms involved, "pre-pulsing" experiments, first described by Sette and co-workers (DeMagistris, M.T. et al. (1992) Cell 68:625-634), were performed. This experimental setting allows agonist (TT830) to bind to MHC Class II in the absence of competition from the inhibitor (TT833S) and thus, only TCR antagonists but not pure MHC

blockers would be effective in inhibiting agonist-stimulated T cell proliferation. Ag-presenting monocytes were pulsed with suboptimal (20 nM) TT830 for 4 h to allow TT830 to bind with MHC Class II in the absence of competition from TT833S. The APL, TT833S or Fab22-TT833S, was then incubated with the pre-pulsed monocytes for an additional 16 h.

5 Responding T cells were added and their proliferation was measured as described above. Even under such conditions where MHC blockade plays a minimal role, T cell proliferation was still inhibited (Figure 33). Thus, the inhibition appears to be a result of competition for T cell receptor rather than for MHC Class II binding.

10 *Presentation of TT833S and Fab22-TT833S fails to stimulate the production of IL-4 and IFN-γ by T cells*

In some circumstances, APL can stimulate T cell cytokine production but not proliferation (Evavold B.D. and Allen P. M. (1991) *Science* 252:1308-1310). To determine if presentation of TT833S simulates the production of cytokines, the level of IL-4 and IFN-γ in supernatants obtained from Ag presentation assays was determined. As shown in Figure 34, both TT833S and Fab22-TT833S were ineffective in stimulating IFN-γ and IL-4 production by T cells.

*Presentation of TT833S and Fab22-TT833S does not lead to T cell anergy*

20 Allen and co-workers reported that interaction of TCR with some APL-MHC Class II complexes led to T cell anergy (Sloan-Lancaster J. et al. (1993) *Nature* 363:156-159, Sloan-Lancaster J. et al. (1994) *J. Exp. Med.* 180:1195-1205). An experimental scheme similar to theirs was used to determine whether presentation of TT833S could also cause T cell anergy. As shown in Figure 35, when T cells were recovered after incubation with APC and TT833S or Fab22-TT833S for 1 day, 2 days, or 4 days, they responded to subsequent antigenic challenge as well as T cells which had been incubated with APC alone. Therefore, the antagonist, presented via the use of either peptide alone or Fab22-TT833S, did not cause T cell anergy. Furthermore, the same percentage of viable T cells (about 50%) was recovered from cultures with no peptides, TT833S or Fab22-TT833S, suggesting that presentation of 25 TT833S also did not increase T cell death.

30 The observation that immunogenicity is increased by about 1000 fold by targeting antigenic peptides to Fc<sub>Y</sub>RI using an anti-Fc<sub>Y</sub>RI mAb 22-based fusion protein indicates that such fusion proteins will be useful for peptide-based vaccines for, e.g., 35 infectious diseases and cancer. Engineering peptides into the constant domains of human mAb that are specific for particular APC surface molecules represent a general approach to increase the antigenic potency for peptide-based vaccines.

Moreover, the observation that the Fc<sub>Y</sub>RI-targeted antagonistic peptide inhibited proliferation of TT830-specific T cells even when APCs were first pulsed with native peptide, a situation comparable to that which would be encountered *in vivo* when attempting to ameliorate an autoimmune response show that targeted presentation of 5 antagonistic peptides can be used as Ag-specific therapies for disorders, e.g., T cell-mediated autoimmune diseases. APL-based treatment will provide an antigen-specific immunotherapy for T cell mediated autoimmune diseases such as rheumatoid arthritis and multiple sclerosis. Furthermore, the use of a fusion protein having one binding specificity to an Fc<sub>Y</sub>RI and a peptide which is a partial agonist of an antigen involved in immune disorders characterized 10 by excessive immune responses will be useful in treating such immune disorders by inducing antigen-specific anergy. Thus, the invention provides methods for treating various immunological disorders, by providing a method allowing for increased antigen presentation of antigens, which either stimulate T cells, block T cell proliferation and/or cytokine secretion, or which induce anergy in the T cells.

15

**Example 8: Functional Single Chain anti-Fc<sub>Y</sub>RI-anti-CEA Bispecific Molecules**

This example demonstrates that a recombinant bispecific single chain 20 molecule comprising a humanized anti-Fc<sub>Y</sub>RI antibody fused to an anti-carcinoembryonic (anti-CEA) antibody is capable of binding to Fc<sub>Y</sub>RI and to CEA.

Figure 37 is a schematic representation of mammalian expression constructs 25 encoding bispecific single chain molecules (constructs 321 and 323) having one binding specificity for the Fc<sub>Y</sub>RI and one binding specificity for carcinoembryonic antigen (CEA) that were prepared. The amino acid sequence of the bispecific single chain molecule H22-anti-CEA encoded by construct 321 (SEQ ID NO: 16) and the nucleic acid encoding this fusion protein (SEQ ID NO: 15) are shown in Figure 40. The bispecific single chain molecule H22-anti-CEA encoded by construct 323 differs from the fusion protein encoded by construct 321. 30 only in that the VH and VL chains of H22 were switched.

A mammalian expression construct encoding a single chain antibody having one binding specificity for the Fc<sub>Y</sub>RI (construct 225) was also prepared. The amino acid sequence of the single chain antibody H22 encoded by construct 225 (SEQ ID NO: 14) and 35 the nucleic acid encoding this single chain antibody (SEQ ID NO: 13) are shown in Figure 39.

Each of these constructs were cloned into the Hind III and XbaI sites of pcDNA3 (InVitrogen), from which expression is driven from the CMV promoter. Each of

these constructs also contain a nucleic acid sequence encoding a peptide from c-myc and a hexa-histidine peptide, which were used for purification of the recombinant protein from the cell culture. The c-myc tag corresponds to amino acids 410 to 420 of human c-myc (Evan et al. (1985) *Mol. Cell. Biol.* 5:3610). The anti-CEA single chain antibody, termed MFE-23, is 5 further described in Casey et al. (1994) *J. Immunol. Methods* 179:105 and Chester et al. (1994) *Lancet* 343:455.

The single chain bispecific molecules H22-anti-CEA and the single chain H22 antibody were used in binding assays, performed as follows. ELISA plates are coated with 10 CEA and the blocked with 5% PBA. Supernatants of the cells transfected with the constructs encoding the single chain molecules (transfectomas) were added to the plates, soluble Fc<sub>y</sub> RI/IgM- $\mu$  (supernatant from COS transfected cells, described above) was added and binding was detected by incubation of the plates with alkaline-phosphatase (AP) conjugated goat anti-human IgM, development with PNPP, and reading of the plate at 405-650 nm.

15

The results are presented in Figure 38. The results indicate that the single chain bispecific H22-anti-CEA molecules encoded by constructs 321 and 323 bind both Fc<sub>y</sub> RI and CEA. On the other hand, the single chain H22 antibody (encoded by construct 225) does not bind both Fc<sub>y</sub> RI and CEA, as expected.

20

#### Equivalents

Those skilled in the art will recognize, or be able to ascertain using no more than routine experimentation, many equivalents to the specific embodiments of the invention 25 described herein. Such equivalents are intended to be encompassed by the following claims.

## SEQUENCE LISTING

## (1) GENERAL INFORMATION:

5

## (i) APPLICANT:

- (A) NAME: Medarex, Inc.
- (B) STREET: 1545 Route 22 East
- (C) CITY: Annandale
- 10 (D) STATE: New Jersey
- (E) COUNTRY: USA
- (F) POSTAL CODE (ZIP): 08801-0953

10

(ii) TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED  
15 OF ANTI-Fc RECEPTOR ANTIBODIES

15

## (iii) NUMBER OF SEQUENCES: 16

20

## (iv) COMPUTER READABLE FORM:

- (A) MEDIUM TYPE: Floppy disk
- (B) COMPUTER: IBM PC compatible
- (C) OPERATING SYSTEM: PC-DOS/MS-DOS
- (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)

25

## (v) CURRENT APPLICATION DATA:

## (vi) PRIOR APPLICATION DATA:

- (A) APPLICATION NUMBER: US 08/484,172
- (B) FILING DATE: 07-JUNE-1995

30

## (vii) CORRESPONDENCE ADDRESS:

- (A) ADDRESSEE: LAHIVE & COCKFIELD
- (B) STREET: 60 State Street, Suite 510
- (C) CITY: Boston
- 35 (D) STATE: Massachusetts
- (E) COUNTRY: USA
- (F) ZIP: 02109-1875

35

## (viii) ATTORNEY/AGENT INFORMATION:

40

- (A) NAME: Arnold, Beth E.
- (B) REGISTRATION NUMBER: 35,430
- (C) REFERENCE/DOCKET NUMBER: MXI-043CPPC

45

## (2) INFORMATION FOR SEQ ID NO:1:

50

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 24 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: cDNA

55

## (ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..24

- 47 -

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

ACT CAC ACA TGC CCA CCG TGC CCA  
Thr His Thr Cys Pro Pro Cys Pro  
5 1 5

24

## (2) INFORMATION FOR SEQ ID NO:2:

## (i) SEQUENCE CHARACTERISTICS:

10 (A) LENGTH: 27 base pairs  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear

15 (iii) MOLECULE TYPE: cDNA

## (ix) FEATURE:

20 (A) NAME/KEY: CDS  
(B) LOCATION: 1..19

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

25 ACT CAC ACA TGC CCA CCG T GAGGATCC  
Thr His Thr Cys Pro Pro  
1 5

27

## (2) INFORMATION FOR SEQ ID NO:3:

30 (i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 42 base pairs  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
35 (D) TOPOLOGY: linear

40 (iii) MOLECULE TYPE: cDNA

45 (ix) FEATURE:  
(A) NAME/KEY: CDS  
(B) LOCATION: 1..34

ACT CAC ACA TGC TCG AGC CTT CAC GGC GGC CGC T GAGGATCC  
Thr His Thr Cys Ser Ser Leu His Gly Gly Arg

42

50 1 5 10

## (2) INFORMATION FOR SEQ ID NO:4:

55 (i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 300 amino acids  
(B) TYPE: amino acid  
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

- 48 -

(v) FRAGMENT TYPE: internal

5 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Glu Val Gln Leu Val Glu Ser Gly Gly Val Val Gln Pro Gly Arg  
1 5 10 15

10 Ser Leu Arg Leu Ser Cys Ser Ser Ser Gly Phe Ile Phe Ser Asp Asn  
20 25 30

Tyr Met Tyr Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
35 40 45

15 Ala Thr Ile Ser Asp Gly Gly Ser Tyr Thr Tyr Tyr Pro Asp Ser Val  
50 55 60

20 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Phe  
65 70 75 80

Leu Gln Met Asp Ser Leu Arg Pro Glu Asp Thr Gly Val Tyr Phe Cys  
85 90 95

25 Ala Arg Gly Tyr Tyr Arg Tyr Glu Gly Ala Met Asp Tyr Trp Gly Gln  
100 105 110

Gly Thr Pro Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val  
115 120 125

30 Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala  
130 135 140

Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Arg Val Thr Val Ser  
35 145 150 155 160

Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val  
165 170 175

40 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro  
180 185 190

Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
195 200 205

45 Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp  
210 215 220

50 Lys Thr His Thr Cys Ser Thr Thr Ser Thr Gly Thr Ser His Leu  
225 230 235 240

Val Lys Cys Ala Glu Lys Glu Lys Thr Phe Cys Val Asn Gly Gly Glu  
245 250 255

55 Cys Phe Met Val Lys Asp Leu Ser Asn Pro Ser Arg Tyr Leu Cys Lys  
260 265 270

Cys Pro Asn Glu Phe Thr Gly Asp Arg Cys Gln Asn Tyr Val Met Ala  
275 280 285

Ser Phe Tyr Lys Ala Glu Glu Leu Tyr Gln Lys Arg  
290 295 300

## 5 (2) INFORMATION FOR SEQ ID NO:5:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 14 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

10 (ii) MOLECULE TYPE: peptide  
15 (v) FRAGMENT TYPE: internal

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

20 Gln Arg Leu Gly Asn Gln Trp Ala Val Gly His Leu Met Gly  
1 5 10

## (2) INFORMATION FOR SEQ ID NO:6:

25 (i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 15 amino acids  
(B) TYPE: amino acid  
(D) TOPOLOGY: linear

30 (ii) MOLECULE TYPE: peptide  
(v) FRAGMENT TYPE: internal

35 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:  
Gln Tyr Ile Lys Ala Asn Ser Lys Phe Ile Gly Ile Thr Glu Leu  
1 5 10 15

## 40 (2) INFORMATION FOR SEQ ID NO:7:

45 (i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 54 base pairs  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear

50 (ii) MOLECULE TYPE: cDNA

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

55 TCGAGCCAGT ACATCAAGGC GAATTCCAAG TTCATCGGCA TCACCGAGCT CTGA

54

## (2) INFORMATION FOR SEQ ID NO:8:

## (i) SEQUENCE CHARACTERISTICS:

- 50 -

- (A) LENGTH: 53 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

5

(ii) MOLECULE TYPE: cDNA

10

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

CGGTCATGTA GTTCCGCTTA AGGTTCAAGT AGCCGTAGTG GCTCGAGACT CCG

53

15

(2) INFORMATION FOR SEQ ID NO:9:

20

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids

- (B) TYPE: amino acid

- (D) TOPOLOGY: linear

25

(ii) MOLECULE TYPE: peptide

(v) FRAGMENT TYPE: internal

25

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

30

Gln Tyr Ile Ser Ala Asn Ser Lys Phe Ile Gly Ile Thr Glu Leu  
1 5 10 15

35

(2) INFORMATION FOR SEQ ID NO:10:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 54 base pairs

- (B) TYPE: nucleic acid

- (C) STRANDEDNESS: single

- (D) TOPOLOGY: linear

40

(ii) MOLECULE TYPE: cDNA

45

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

TCGAGCCAGT ACATCAGCGC GAATTCCAAG TTCATCGGCA TCACCGAGCT CTGA

54

50

(2) INFORMATION FOR SEQ ID NO:11:

55

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 53 base pairs

- (B) TYPE: nucleic acid

- (C) STRANDEDNESS: single

- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

- 51 -

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

CGGTCATGTA GTCGCGCTTA AGGTTCAAGT AGCCGTAGTG GCTCGAGACT CCG

53

## 5 (2) INFORMATION FOR SEQ ID NO:12:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 21 amino acids
- (B) TYPE: amino acid
- 10 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(v) FRAGMENT TYPE: internal

15

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

20 Phe Asn Asn Phe Thr Val Ser Phe Trp Leu Arg Val Pro Lys Val Ser  
1 5 10 15Ala Ser His Leu Glu  
20

25

## (2) INFORMATION FOR SEQ ID NO:13:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 913 base pairs
- 30 (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

35

## (ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 11..911

40

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

45 AAGCTTCACC ATG GGA TGG AGC TGT ATC ATC CTC TTC TTG GTG GCC ACA  
Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr  
1 5 10

49

50 GCT ACC GGT GTC CAC TCC GAT ATC CAA CTG GTG GAG AGC GGT GGA GGT  
Ala Thr Gly Val His Ser Asp Ile Gln Leu Val Glu Ser Gly Gly  
15 20 25

97

55 GTT GTG CAA CCT GGC CGG TCC CTG CGC CTG TCC TGC TCC TCG TCT GGC  
Val Val Gln Pro Gly Arg Ser Leu Arg Leu Ser Cys Ser Ser Gly  
30 35 40 45

145

TTC AGT TTC AGT GAC AAT TAC ATG TAT TGG GTG AGA CAG GCA CCT GGA  
Phe Ile Phe Ser Asp Asn Tyr Met Tyr Trp Val Arg Gln Ala Pro Gly  
50 55 60

193

- 52 -

	AAA GGT CTT GAG TGG GTT GCA ACC ATT AGT GAT GGT GGT AGT TAC ACC Lys Gly Leu Glu Trp Val Ala Thr Ile Ser Asp Gly Ser Tyr Thr 65 70 75	241
5	TAC TAT CCA GAC AGT GTG AAG GGA AGA TTT ACA ATA TCG AGA GAC AAC Tyr Tyr Pro Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn 80 85 90	289
10	AGC AAG AAC ACA TTG TTC CTG CAA ATG GAC AGC CTG AGA CCC GAA GAC Ser Lys Asn Thr Leu Phe Leu Gln Met Asp Ser Leu Arg Pro Glu Asp 95 100 105	337
15	ACC GGG GTC TAT TTT TGT GCA AGA GGC TAC TAT AGG TAC GAG GGG GCT Thr Gly Val Tyr Phe Cys Ala Arg Gly Tyr Tyr Arg Tyr Glu Gly Ala 110 115 120 125	385
	ATG GAC TAC TGG GGC CAA GGG ACC CCG GTC ACC GTG AGC TCA GGA GGT Met Asp Tyr Trp Gly Gln Gly Thr Pro Val Thr Val Ser Ser Gly Gly 130 135 140	433
20	GGC GGC TCC GGA GGT GGA GGC AGC GGA GGG GGC GGA TCC GAC ATC CAG Gly Gly Ser Gly Gly Ser Gly Gly Ser Gly Ser Asp Ile Gln 145 150 155	481
25	CTG ACC CAG AGC CCA AGC AGC CTG AGC GCC AGC GTG GGT GAC AGA GTG Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp Arg Val 160 165 170	529
30	ACC ATC ACC TGT AAG TCC AGT CAA AGT GTT TTA TAC AGT TCA AAT CAG Thr Ile Thr Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser Ser Asn Gln 175 180 185	577
35	AAG AAC TAC TTG GCC TGG TAC CAG CAG AAG CCA GGT AAG GCT CCA AAG Lys Asn Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys 190 195 200 205	625
	CTG CTG ATC TAC TGG GCA TCC ACT AGG GAA TCT GGT GTG CCA AGC AGA Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val Pro Ser Arg 210 215 220	673
40	TTC AGC GGT AGC GGT AGC GGT ACC GAC TTC ACC TTC ACC ATC AGC AGC Phe Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser 225 230 235	721
45	CTC CAG CCA GAG GAC ATC GCC ACC TAC TAC TGC CAT CAA TAC CTC TCC Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys His Gln Tyr Leu Ser 240 245 250	769
50	TCG TGG ACG TTC GGC CAA GGG ACC AAG GTG GAA ATC AAA TCT AGC TGC Ser Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Ser Ser Cys 255 260 265	817
55	TCG AGC GGA GGC GGG GGT AGC GAT ATC GCG GCC GCA GAA CAG AAA CTC Ser Ser Gly Gly Ser Asp Ile Ala Ala Ala Glu Gln Lys Leu 270 275 280 285	865
	ATC TCA GAA GAG GAT CTG AAT GGC GCC GCA CAT CAC CAT CAT CAC CAT Ile Ser Glu Glu Asp Leu Asn Gly Ala Ala His His His His His His 290 295 300	913

## (2) INFORMATION FOR SEQ ID NO:14:

## (i) SEQUENCE CHARACTERISTICS:

5 (A) LENGTH: 301 amino acids  
 (B) TYPE: amino acid  
 (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: protein

## 10 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly  
 1 5 10 15

15 Val His Ser Asp Ile Gln Leu Val Glu Ser Gly Gly Val Val Gln  
 20 25 30

20 Pro Gly Arg Ser Leu Arg Leu Ser Cys Ser Ser Ser Gly Phe Ile Phe  
 35 40 45

Ser Asp Asn Tyr Met Tyr Trp Val Arg Gln Ala Pro Gly Lys Gly Leu  
 50 55 60

25 Glu Trp Val Ala Thr Ile Ser Asp Gly Gly Ser Tyr Thr Tyr Tyr Pro  
 65 70 75 80

Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn  
 85 90 95

30 Thr Leu Phe Leu Gln Met Asp Ser Leu Arg Pro Glu Asp Thr Gly Val  
 100 105 110

35 Tyr Phe Cys Ala Arg Gly Tyr Tyr Arg Tyr Glu Gly Ala Met Asp Tyr  
 115 120 125

Trp Gly Gln Gly Thr Pro Val Thr Val Ser Ser Gly Gly Gly Ser  
 130 135 140

40 Gly Gly Gly Ser Gly Gly Ser Asp Ile Gln Leu Thr Gln  
 145 150 155 160

Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr  
 165 170 175

45 Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser Ser Asn Gln Lys Asn Tyr  
 180 185 190

50 Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
 195 200 205

Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val Pro Ser Arg Phe Ser Gly  
 210 215 220

55 Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Leu Gln Pro  
 225 230 235 240

Glu Asp Ile Ala Thr Tyr Tyr Cys His Gln Tyr Leu Ser Ser Trp Thr  
 245 250 255

Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Ser Ser Cys Ser Ser Gly  
 260 265 270

5 Gly Gly Gly Ser Asp Ile Ala Ala Ala Glu Gln Lys Leu Ile Ser Glu  
 275 280 285

Glu Asp Leu Asn Gly Ala Ala His His His His His His  
 290 295 300

10

(2) INFORMATION FOR SEQ ID NO:15:

## (i) SEQUENCE CHARACTERISTICS:

15

- (A) LENGTH: 1679 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

20

## (ii) MOLECULE TYPE: cDNA

25

## (ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 11..1667

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

30	AAGCTTCACC ATG GGA TGG AGC TGT ATC ATC CTC TTC TTG GTG GCC ACA Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr 1 5 10	49
35	GCT ACC GGT GTC CAC TCC GAT ATC CAA CTG GTG GAG AGC GGT GGA GGT Ala Thr Gly Val His Ser Asp Ile Gln Leu Val Glu Ser Gly Gly 15 20 25	97
40	GTT GTG CAA CCT GGC CGG TCC CTG CGC CTG TCC TGC TCC TCG TCT GGC Val Val Gln Pro Gly Arg Ser Leu Arg Leu Ser Cys Ser Ser Ser Gly 30 35 40 45	145
45	TTC ATT TTC AGT GAC AAT TAC ATG TAT TGG GTG AGA CAG GCA CCT GGA Phe Ile Phe Ser Asp Asn Tyr Met Tyr Trp Val Arg Gln Ala Pro Gly 50 55 60	193
50	AAA GGT CTT GAG TGG GTT GCA ACC ATT AGT GAT GGT GGT AGT TAC ACC Lys Gly Leu Glu Trp Val Ala Thr Ile Ser Asp Gly Gly Ser Tyr Thr 65 70 75	241
55	TAC TAT CCA GAC AGT GTG AAG GGA AGA TTT ACA ATA TCG AGA GAC AAC Tyr Tyr Pro Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn 80 85 90	289
60	AGC AAG AAC ACA TTG TTC CTG CAA ATG GAC AGC CTG AGA CCC GAA GAC Ser Lys Asn Thr Leu Phe Leu Gln Met Asp Ser Leu Arg Pro Glu Asp 95 100 105	337
65	ACC GGG GTC TAT TTT TGT GCA AGA GGC TAC TAT AGG TAC GAG GGG GCT Thr Gly Val Tyr Phe Cys Ala Arg Gly Tyr Tyr Arg Tyr Glu Gly Ala 110 115 120 125	385

	ATG GAC TAC TGG GGC CAA GGG ACC CCG GTC ACC GTG AGC TCA GGA GGT Met Asp Tyr Trp Gly Gln Gly Thr Pro Val Thr Val Ser Ser Gly Gly 130 135 140	433
5	GGC GGC TCC GGA GGT GGA GGC AGC GGA GGG GGC GGA TCC GAC ATC CAG Gly Gly Ser Gly Gly Ser Gly Gly Ser Gly Gly Ser Asp Ile Gln 145 150 155	481
10	CTG ACC CAG AGC CCA AGC AGC CTG AGC GCC AGC GTG GGT GAC AGA GTG Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp Arg Val 160 165 170	529
15	ACC ATC ACC TGT AAG TCC AGT CAA AGT GTT TTA TAC AGT TCA AAT CAG Thr Ile Thr Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser Ser Asn Gln 175 180 185	577
20	AAG AAC TAC TTG GCC TGG TAC CAG CAG AAG CCA GGT AAG GCT CCA AAG Lys Asn Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys 190 195 200 205	625
25	CTG CTG ATC TAC TGG GCA TCC ACT AGG GAA TCT GGT GTG CCA AGC AGA Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val Pro Ser Arg 210 215 220	673
30	TTC AGC GGT AGC GGT AGC GGT ACC GAC TTC ACC TTC ACC ATC AGC AGC Phe Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser 225 230 235	721
35	TCG TGG ACG TTC GGC CAA GGG ACC AAG GTG GAA ATC AAA TCT AGC TGC Ser Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Ser Ser Cys 255 260 265	817
40	TCG AGC GGA GGG GGG GGT AGC GAT ATC AAA CTG CAG CAG TCT GGG GCA Ser Ser Gly Gly Ser Asp Ile Lys Leu Gln Gln Ser Gly Ala 270 275 280 285	865
45	GAA CTT GTG AGG TCA GGG ACC TCA GTC AAG TTG TCC TGC ACA GCT TCT Glu Leu Val Arg Ser Gly Thr Ser Val Lys Leu Ser Cys Thr Ala Ser 290 295 300	913
50	GGC TTC AAC ATT AAA GAC TCC TAT ATG CAC TGG TTG AGG CAG GGG CCT Gly Phe Asn Ile Lys Asp Ser Tyr Met His Trp Leu Arg Gln Gly Pro 305 310 315	961
55	GAA CAG GGC CTG GAG TGG ATT GGA TGG ATT GAT CCT GAG AAT GGT GAT Glu Gln Gly Leu Glu Trp Ile Gly Trp Ile Asp Pro Glu Asn Gly Asp 320 325 330	1009
	ACT GAA TAT GCC CCG AAG TTC CAG GGC AAG GCC ACT TTT ACT ACA GAC Thr Glu Tyr Ala Pro Lys Phe Gln Gly Lys Ala Thr Phe Thr Thr Asp 335 340 345	1057
	ACA TCC TCC AAC ACA GCC TAC CTG CAG CTG AGC AGC CTG ACA TCT GAG Thr Ser Ser Asn Thr Ala Tyr Leu Gln Leu Ser Ser Leu Thr Ser Glu	1105

- 56 -

	350	355	360	365	
5	GAC ACT GCC GTC TAT TAT TGT AAT GAG GGG ACT CCG ACT GGG CCG TAC Asp Thr Ala Val Tyr Tyr Cys Asn Glu Gly Thr Pro Thr Gly Pro Tyr				1153
	370	375	380		
10	TAC TTT GAC TAC TGG GGC CAA GGG ACC ACG GTC ACC GTC TCC TCA GGT Tyr Phe Asp Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly				1201
	385	390	395		
15	GGA GGC GGT TCA GGC GGA GGT GGC TCT GGC GGT GGC GGA TCA GAA AAT Gly Gly Ser Gly Gly Ser Gly Gly Ser Gly Gly Ser Gly Asn				1249
	400	405	410		
20	GTG CTC ACC CAG TCT CCA GCA ATC ATG TCT GCA TCT CCA GGG GAG AAG Val Leu Thr Gln Ser Pro Ala Ile Met Ser Ala Ser Pro Gly Glu Lys				1297
	415	420	425		
25	GTC ACC ATA ACC TGC AGT GCC AGC TCA AGT GTA AGT TAC ATG CAC TGG Val Thr Ile Thr Cys Ser Ala Ser Ser Val Ser Tyr Met His Trp				1345
	430	435	440	445	
	TTC CAG CAG AAG CCA GGC ACT TCT CCC AAA CTC TGG ATT TAT AGC ACA Phe Gln Gln Lys Pro Gly Thr Ser Pro Lys Leu Trp Ile Tyr Ser Thr				1393
	450	455	460		
30	TCC AAC CTG GCT TCT GGA GTC CCT GCT CGC TTC AGT GGC AGT GGA TCT Ser Asn Leu Ala Ser Gly Val Pro Ala Arg Phe Ser Gly Ser Gly Ser				1441
	465	470	475		
35	GGG ACC TCT TAC TCT CTC ACA ATC AGC CGA ATG GAG GCT GAA GAT GCT Gly Thr Ser Tyr Ser Leu Thr Ile Ser Arg Met Glu Ala Glu Asp Ala				1489
	480	485	490		
40	GCC ACT TAT TAC TGC CAG CAA CGG AGT AGT TAC CCA CTC ACG TTC GGT Ala Thr Tyr Tyr Cys Gln Gln Arg Ser Ser Tyr Pro Leu Thr Phe Gly				1537
	495	500	505		
45	GCT GGC ACC AAG CTG GAG CTG AAA CGG GCG GCA GGC TCG AGC GGA GGC Ala Gly Thr Lys Leu Glu Leu Lys Arg Ala Ala Gly Ser Ser Gly Gly				1585
	510	515	520	525	
	GGG GGT AGC GAT ATC GCG GCC GCA GAA CAG AAA CTC ATC TCA GAA GAG Gly Gly Ser Asp Ile Ala Ala Ala Glu Gln Lys Leu Ile Ser Glu Glu				1633
	530	535	540		
50	GAT CTG AAT GGC GCC GCA CAT CAC CAT CAT CAC CAT TGATTCTAGA Asp Leu Asn Gly Ala Ala His His His His His His				1679
	545	550			

## (2) INFORMATION FOR SEQ ID NO:16:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 553 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: protein

- 57 -

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly  
1 5 10 15

5 Val His Ser Asp Ile Gln Leu Val Glu Ser Gly Gly Val Val Gln  
20 25 30

10 Pro Gly Arg Ser Leu Arg Leu Ser Cys Ser Ser Ser Gly Phe Ile Phe  
35 40 45

15 Ser Asp Asn Tyr Met Tyr Trp Val Arg Gln Ala Pro Gly Lys Gly Leu  
50 55 60

20 15 Glu Trp Val Ala Thr Ile Ser Asp Gly Gly Ser Tyr Thr Tyr Tyr Pro  
65 70 75 80

Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn  
85 90 95

25 25 Tyr Phe Cys Ala Arg Gly Tyr Tyr Arg Tyr Glu Gly Ala Met Asp Tyr  
115 120 125

30 Trp Gly Gln Gly Thr Pro Val Thr Val Ser Ser Gly Gly Gly Ser  
130 135 140

35 30 Gly Gly Gly Ser Gly Gly Ser Asp Ile Gln Leu Thr Gln  
145 150 155 160

Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp Arg Val Thr Ile Thr  
165 170 175

40 35 Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser Ser Asn Gln Lys Asn Tyr  
180 185 190

Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
195 200 205

45 Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val Pro Ser Arg Phe Ser Gly  
210 215 220

50 45 Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Leu Gln Pro  
225 230 235 240

Glu Asp Ile Ala Thr Tyr Tyr Cys His Gln Tyr Leu Ser Ser Trp Thr  
245 250 255

55 50 Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Ser Ser Cys Ser Ser Gly  
260 265 270

Gly Gly Gly Ser Asp Ile Lys Leu Gln Gln Ser Gly Ala Glu Leu Val  
275 280 285

Arg Ser Gly Thr Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn  
290 295 300

- 58 -

Ile Lys Asp Ser Tyr Met His Trp Leu Arg Gln Gly Pro Glu Gln Gly  
305 310 315 320

5 Leu Glu Trp Ile Gly Trp Ile Asp Pro Glu Asn Gly Asp Thr Glu Tyr  
325 330 335

Ala Pro Lys Phe Gln Gly Lys Ala Thr Phe Thr Thr Asp Thr Ser Ser  
340 345 350

10 Asn Thr Ala Tyr Leu Gln Leu Ser Ser Leu Thr Ser Glu Asp Thr Ala  
355 360 365

Val Tyr Tyr Cys Asn Glu Gly Thr Pro Thr Gly Pro Tyr Tyr Phe Asp  
370 375 380

15 Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly  
385 390 395 400

Ser Gly Gly Gly Ser Gly Gly Gly Ser Glu Asn Val Leu Thr  
20 405 410 415

Gln Ser Pro Ala Ile Met Ser Ala Ser Pro Gly Glu Lys Val Thr Ile  
420 425 430

25 Thr Cys Ser Ala Ser Ser Val Ser Tyr Met His Trp Phe Gln Gln  
435 440 445

Lys Pro Gly Thr Ser Pro Lys Leu Trp Ile Tyr Ser Thr Ser Asn Leu  
30 450 455 460

Ala Ser Gly Val Pro Ala Arg Phe Ser Gly Ser Gly Thr Ser  
465 470 475 480

Tyr Ser Leu Thr Ile Ser Arg Met Glu Ala Glu Asp Ala Ala Thr Tyr  
35 485 490 495

Tyr Cys Gln Gln Arg Ser Ser Tyr Pro Leu Thr Phe Gly Ala Gly Thr  
500 505 510

40 Lys Leu Glu Leu Lys Arg Ala Ala Gly Ser Ser Gly Gly Gly Ser  
515 520 525

Asp Ile Ala Ala Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn  
45 530 535 540

Gly Ala Ala His His His His His His  
545 550

*We Claim:*

1. A recombinant multispecific molecule comprising an anti-Fc receptor portion and an anti-target portion.
- 5 2. A recombinant multispecific molecule of claim 1, wherein at least one of the anti-Fc receptor portion or the anti-target portion is humanized.
- 10 3. A recombinant multispecific molecule of claim 2, wherein the anti-Fc receptor portion is an antibody fragment.
4. A recombinant multispecific molecule of claim 2, wherein the anti-target portion is an antibody fragment or a ligand.
- 15 5. A recombinant multispecific molecule of claim 2, wherein the anti-Fc receptor portion binds an Fc receptor on an effector cell at a site which is not bound by endogenous immunoglobulin.
- 20 6. A recombinant multispecific molecule of claim 2, wherein the anti-Fc receptor portion binds an Fc $\gamma$  receptor.
7. A recombinant multispecific molecule of claim 6, wherein the Fc $\gamma$  receptor is a Fc $\gamma$  RI receptor.
- 25 8. A recombinant multispecific molecule of claim 4, wherein the target is a cancer cell.
9. A recombinant multispecific molecule of claim 4, wherein the target is an infectious agent.
- 30 10. A recombinant multispecific molecule of claim 4, wherein the target is an antibody-producing cell.
- 35 11. A recombinant multispecific molecule of claim 8, wherein the target is a breast or ovarian cancer cell.
12. A recombinant multispecific molecule of claim 11, wherein the target is a HER 2/neu expressing cell.

13. A recombinant multispecific molecule of claim 12, wherein the anti-target portion is antibody 520C9.

5 14. A recombinant multispecific molecule of 4, wherein the ligand is epidermal growth factor

15. A multivalent molecule comprising 1) at least one anti-Fc receptor portion, and 2) at least one anti-target portion.

10 16. A multispecific molecule having one anti-FcR, one anti-target portion and one anti-enhancement factor portion.

15 17. The multivalent molecule of claim 15 or 16, wherein the anti-target portion binds a cancer cell.

18. The multivalent molecule of claim 15 or 16, wherein the anti-target portion binds a carcinoma.

20 19. The multivalent molecule of claim 15 or 16, wherein the anti-target portion binds a sarcoma.

25 20. The multivalent molecule of claim 15 or 16, wherein the anti-target portion binds a pathogen.

21. The multivalent molecule of claim 15 or 16, wherein the anti-target portion binds an FcR at an epitope different and distinct from the first FcR monoclonal antibody.

30 22. The multivalent molecule of claim 15 or 16, wherein the anti-target portion binds a soluble protein/peptide.

23. The multivalent molecule of claim 15 or 16, wherein the anti-target portion binds any molecule capable of generating an immune response or monoclonal antibody.

35 24. The multivalent molecule of claim 15 or 16, wherein the binding of the anti-FcR portion to the Fc receptor is not blocked by human immunoglobulin G

25. A multivalent molecule of claim 15 or 16 wherein the anti-FcR portion binds specifically to FcR.

5      26. The multivalent molecule of claim 16, wherein the anti-EF portion binds specifically to a T cell surface antigen.

10      27. The multivalent molecule of claim 16, wherein the anti-EF portion binds CD3.

15      28. The multivalent molecule of claim 16, wherein the anti-EF portion binds a second epitope on FcR.

20      29. The multivalent molecule of claim 16, wherein the anti-EF portion binds a target cell.

15      30. The multivalent molecule of claim 16, wherein the anti-EF portion binds a second FcR.

20      31. A multispecific molecule of claim 16, wherein the anti-EF portion binds specifically to a myeloid-associated cytotoxic trigger molecule.

25      32. A multispecific molecule having one portion that binds specifically to FcγRI, one portion that binds specifically to one epitope of the target antigen and one portion that binds specifically to a second site on the same target cell.

30      33. A method of treating cancer comprising administration of a therapeutically effective amount of a multivalent, multispecific molecule.

30      34. A method of treating autoimmune disease comprising administration of a therapeutically effective amount of a multivalent, multispecific molecule.

35. A method of treating removing unwanted pathogens comprising administration of a therapeutically effective amount of a multivalent, multispecific molecule.

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**b22 hinge region:****[A]**

-ACT CAC ACA TGC CCA CCG TGC CCA — CH<sub>2</sub> — CH<sub>3</sub>  
T H T C P P C P

↓

**[B]**

-ACT CAC ACA TGC CCA CCG TGA GGATCC  
T H T C P P

↓

**[C]**

-ACT CAC ACA TGC XbaI TCG AGC NdeI CIT CAC BamHI GGC GGC CGC TGA GGA TCC  
T H T C S S L H G G R

FIGURE 1

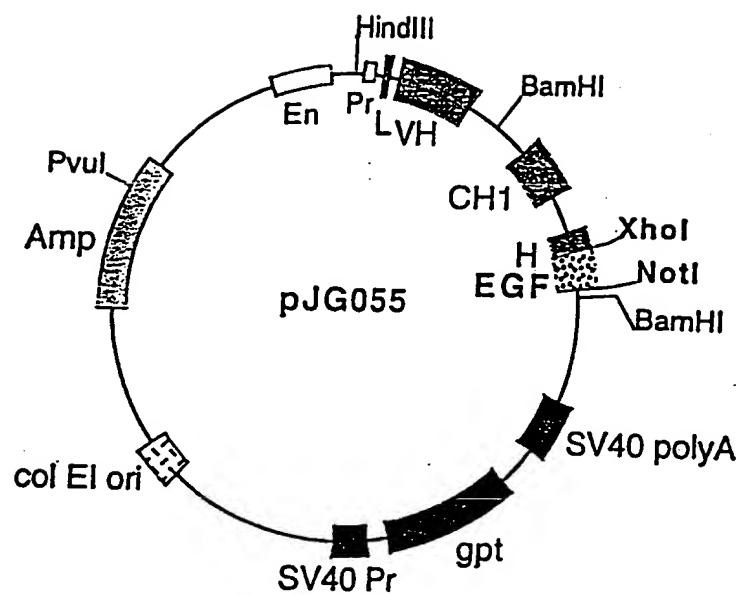


FIGURE 2

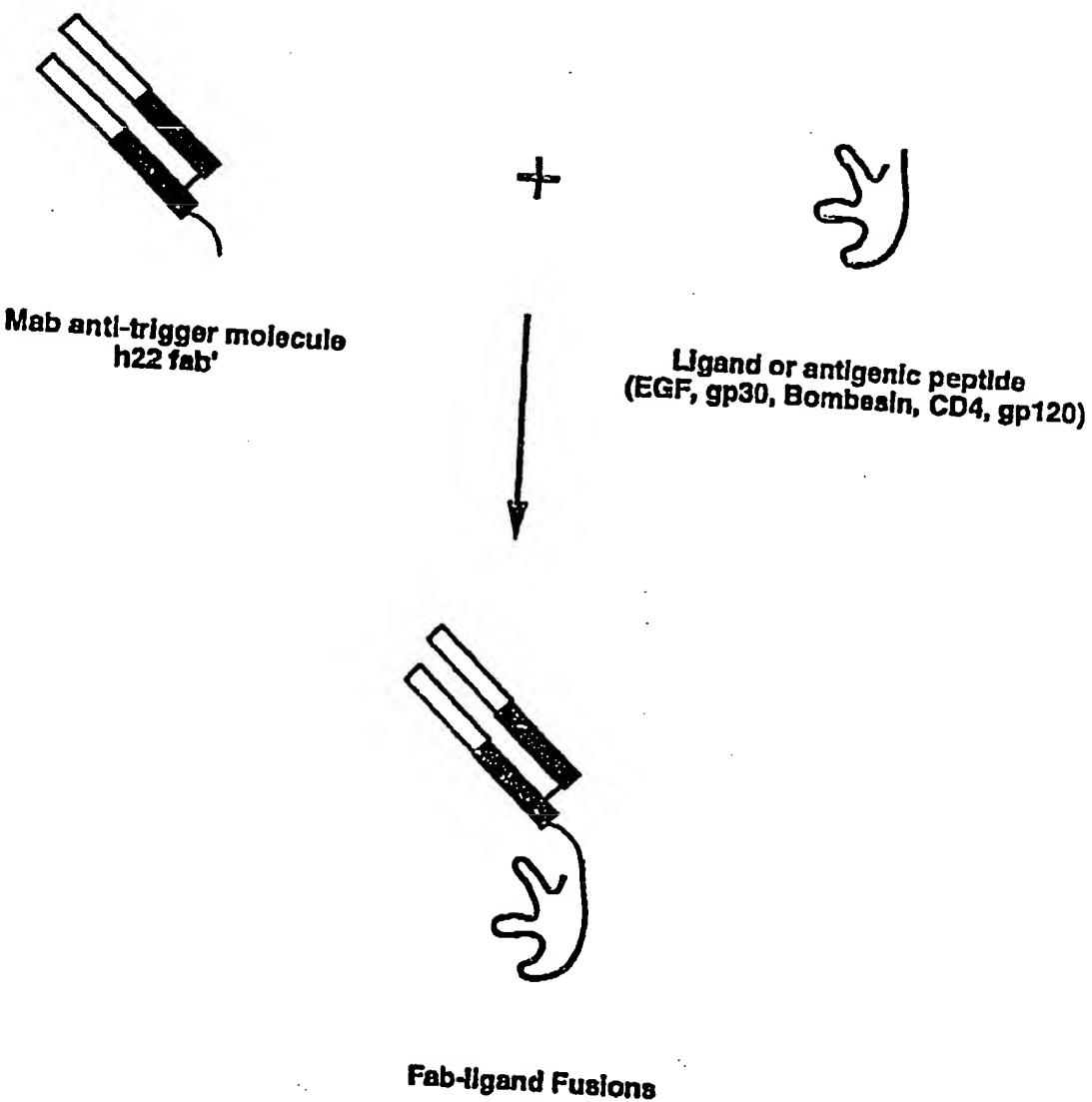


FIGURE 3

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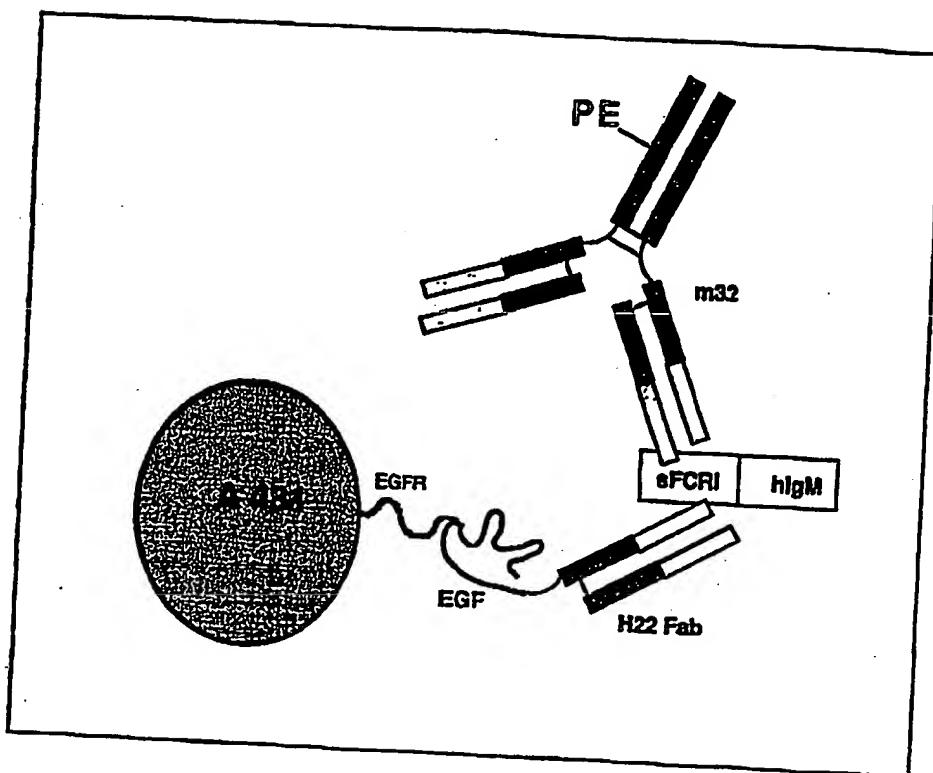


FIGURE 4

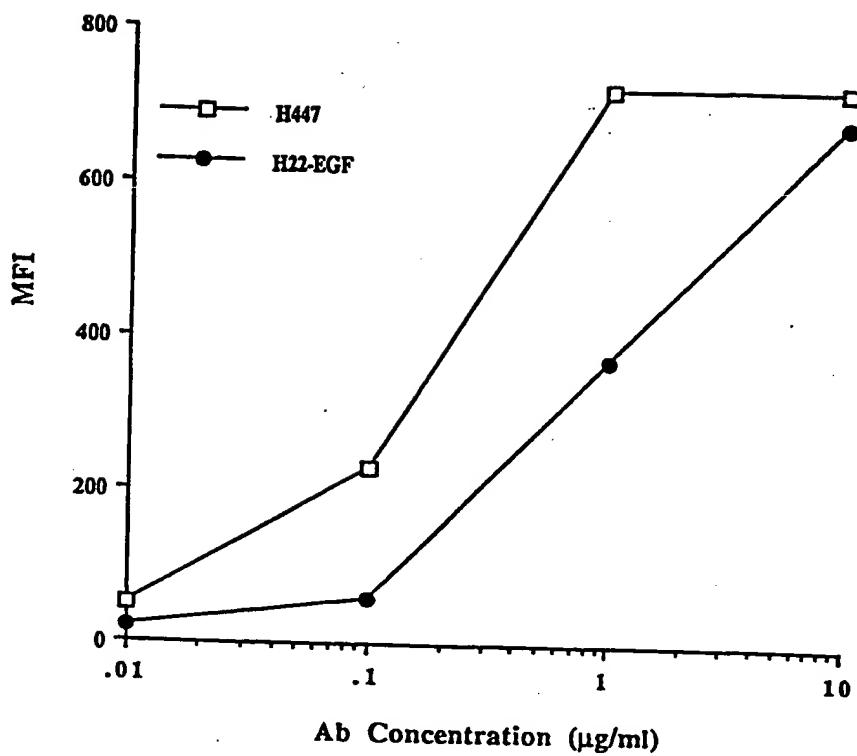


FIGURE 5

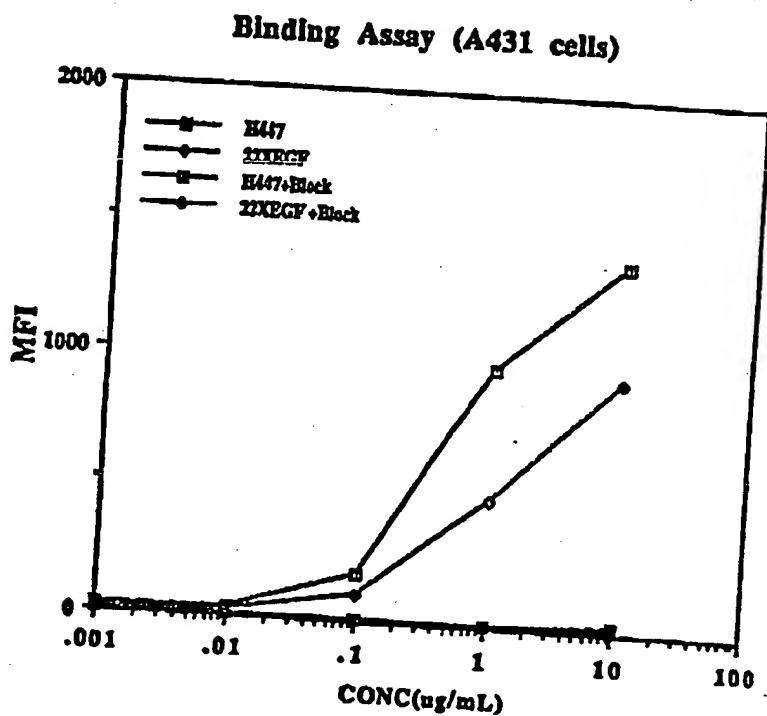


FIGURE 6

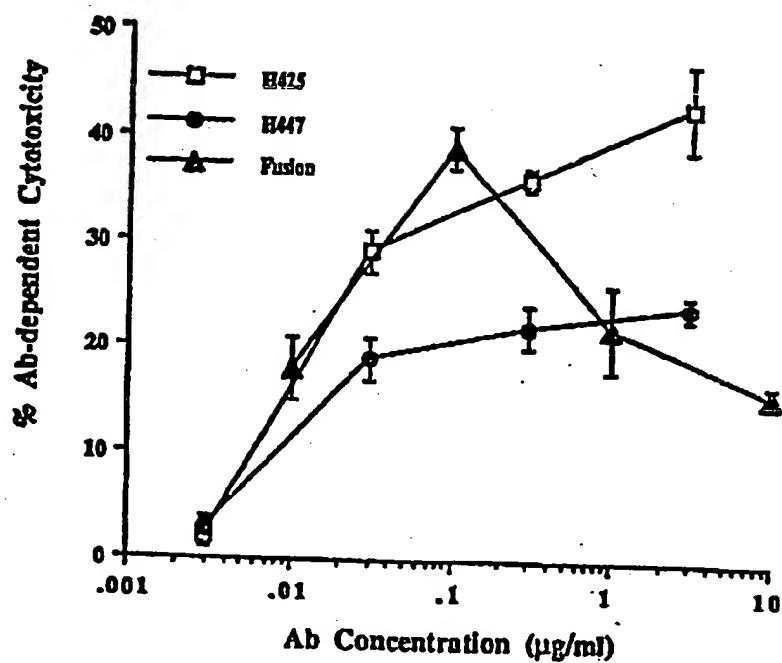


FIGURE 7

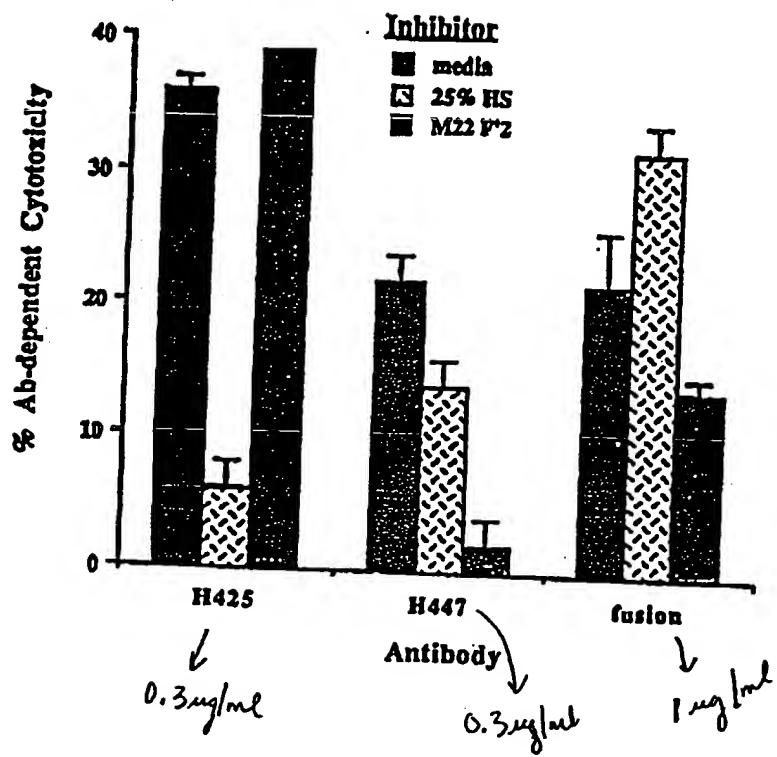


FIGURE 8

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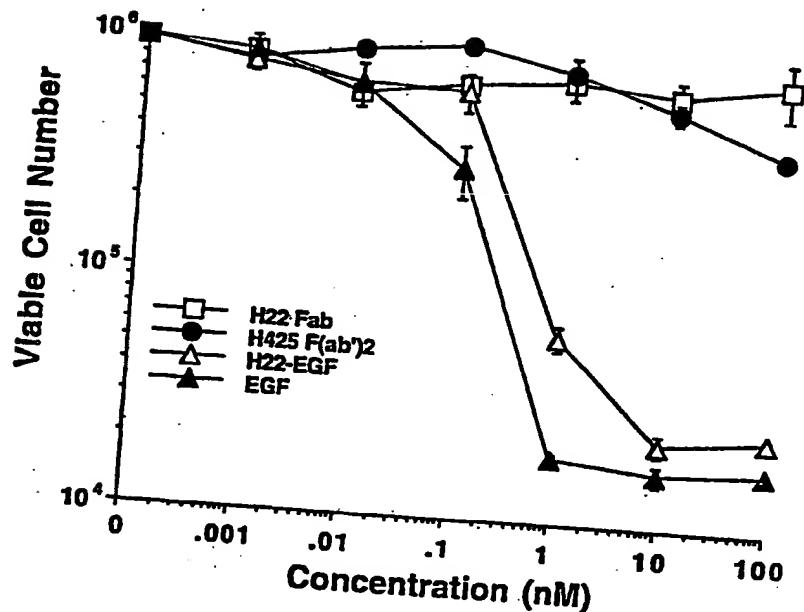


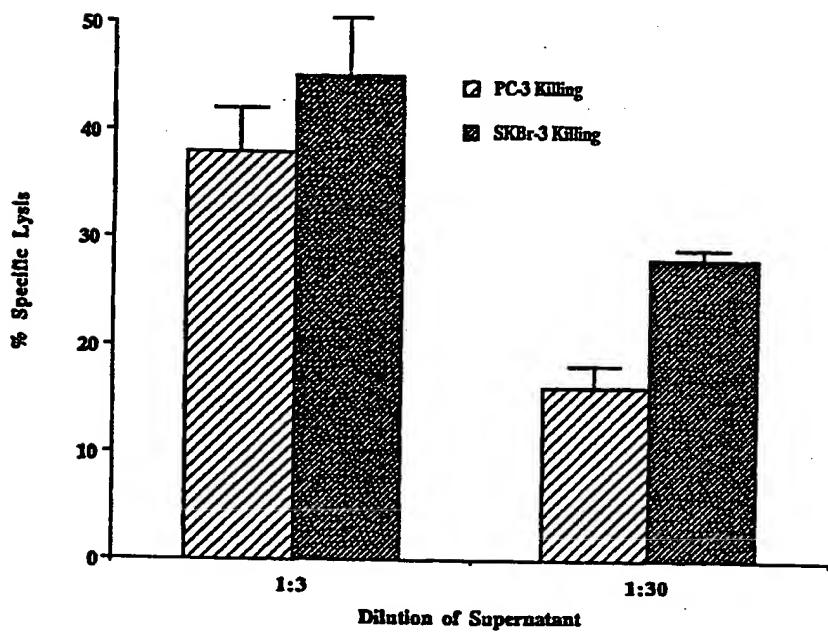
FIGURE 9

10 20 30 40 50  
\* \* \* \* \*  
EVQLVESGGG VVQPGRSLRL SCSSSGPIPS DNMYMYNVRQA PGKGLEWVAT  
60 70 80 90 100  
\* \* \* \* \*  
ISDGGSYTYY PDSVKGRFTI SRDN SKNTLF LQMDSLRPED TGVYFCARGY  
110 120 130 140 150  
\* \* \* \* \*  
YRYEGAMDYW GQGTPVTVSS ASTKGPSVFP LAPSSKSTSG GTAALGCLVK  
160 170 180 190 200  
\* \* \* \* \*  
DYFPERVTVS WNSGALTSGV HTFPAVLQSS GLYSLSSVVT VPSSSLGTQT  
210 220 230 240 250  
\* \* \* \* \*  
YICNVNHRKPS NTKVDKKVEP KSCDKTHTC S TTSTTGTSHL VKCAEKERTF  
H22 Fd Herogulin B2 EGF domain  
260 270 280 290 300  
\* \* \* \* \*  
CVNGGECPMV KDLSNPSRYL CRCPNEFTGD RCQNYVMASP YKAEEELYQKR

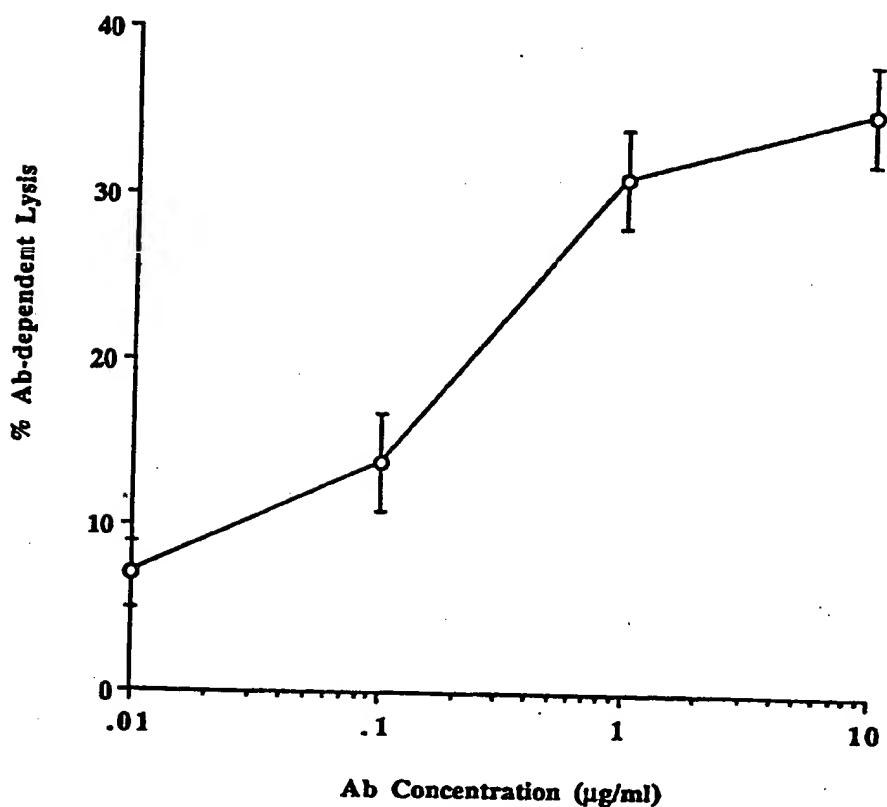
SEQ ID NO: 4

FIGURE 10

**H22-Heregulin Mediated Killing of HER2/3/4-Expressing Tumor Cells  
(IFN-treated Monocytes, E:T=100:1)**



**FIGURE 11**

**H22-Bombesin-Mediated Killing of PC-3 Cells****FIGURE 12**

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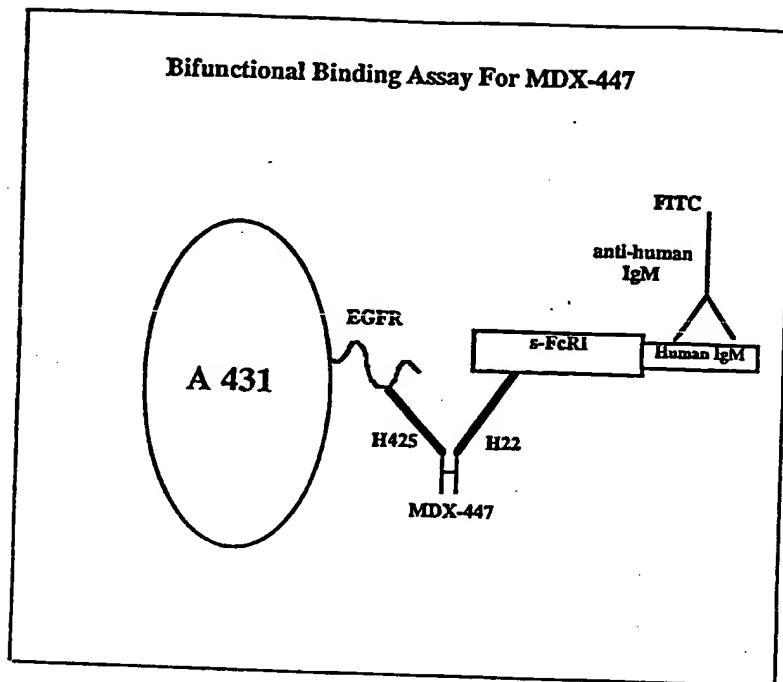


FIGURE 13

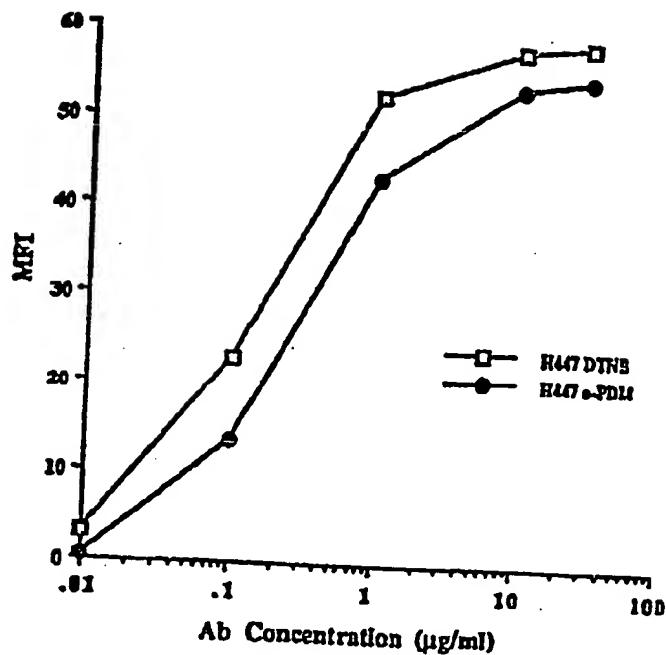


FIGURE 14

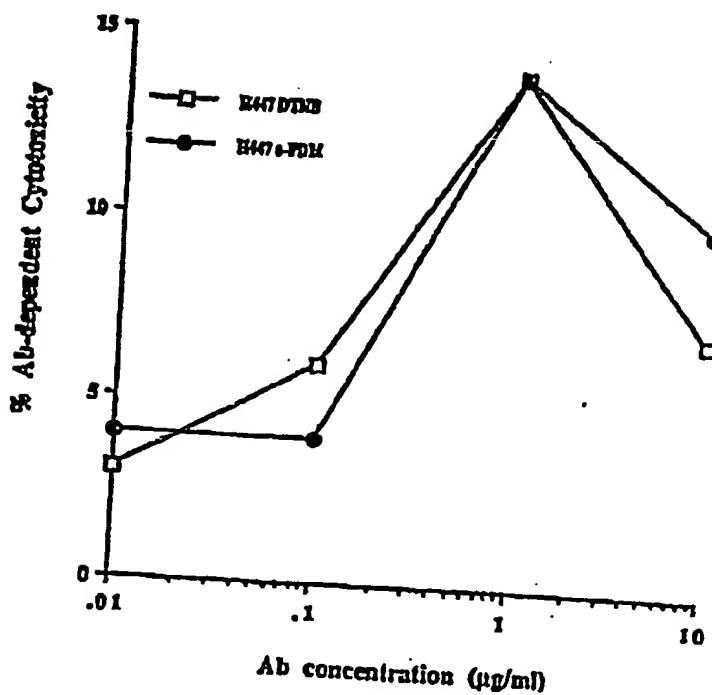


FIGURE 15

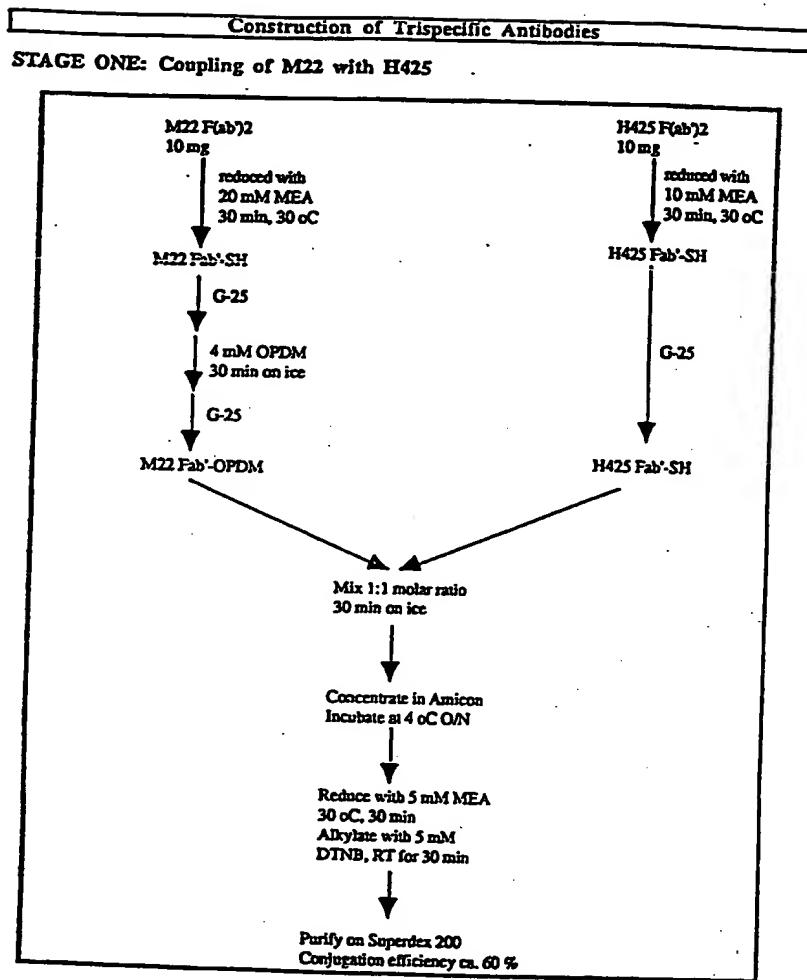


FIGURE 16A

## Construction of Trispecific Antibodies (continued)

## STAGE TWO: Coupling of 520C9 with bispecific M22xH425

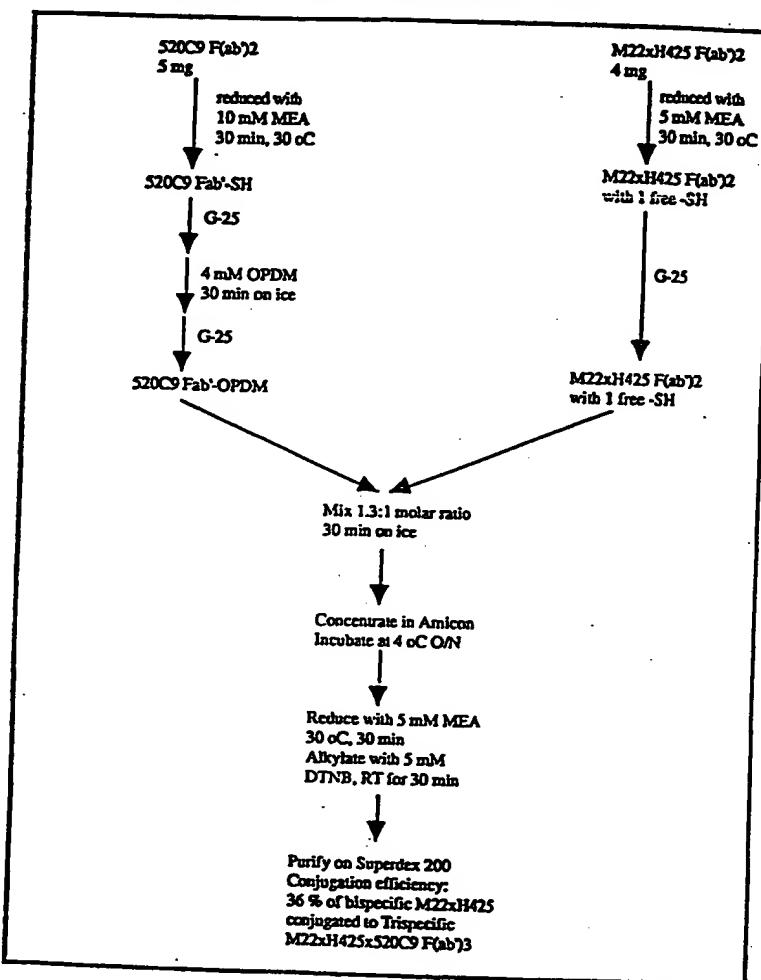


FIGURE 16B

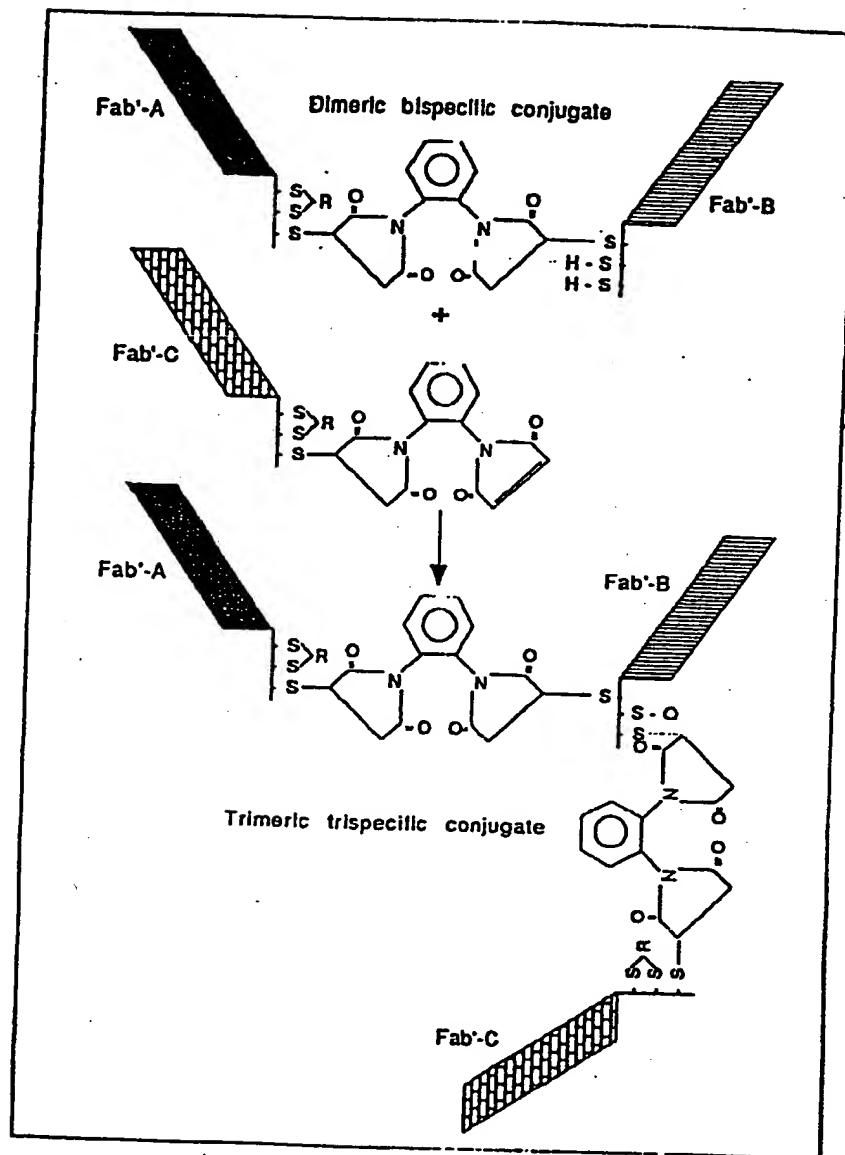


FIGURE 17

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## BIFUNCTIONAL FACS

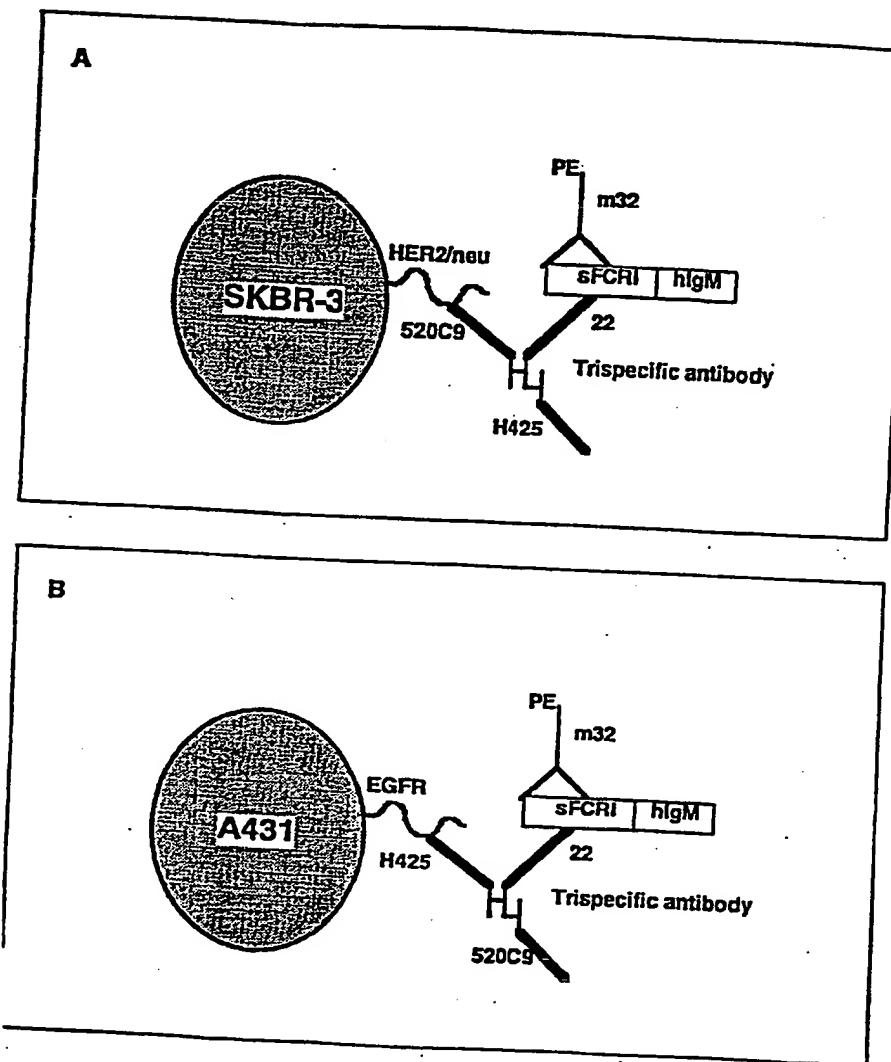


FIGURE 18

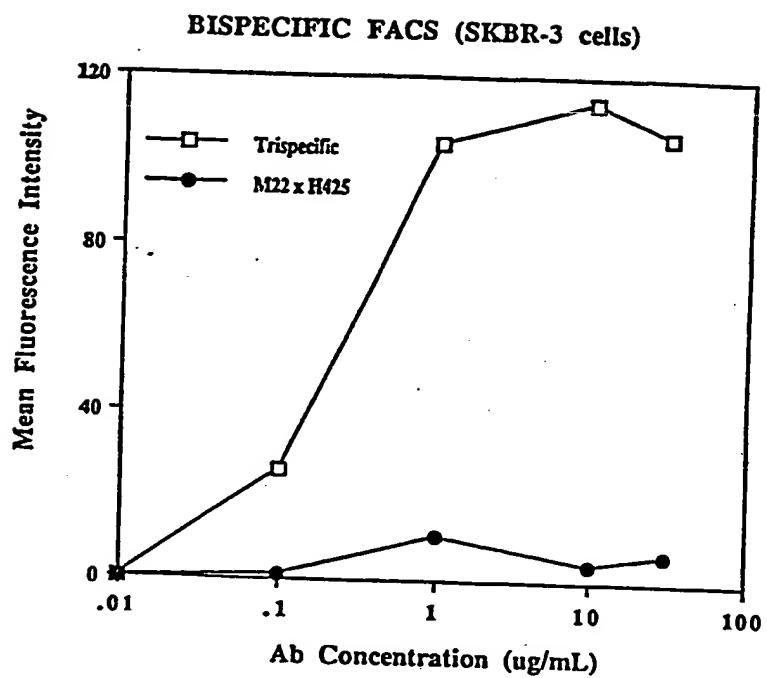


FIGURE 19

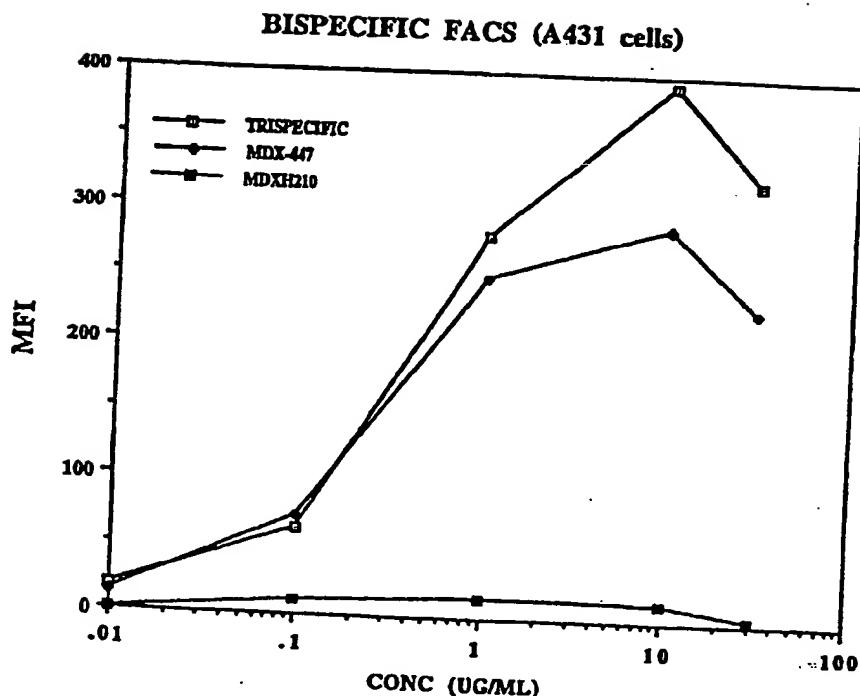


FIGURE 20

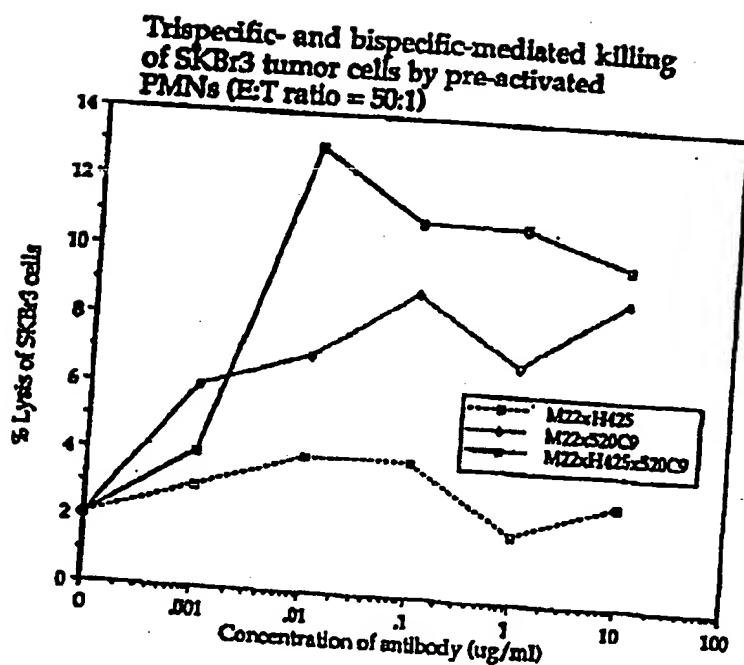


FIGURE 21

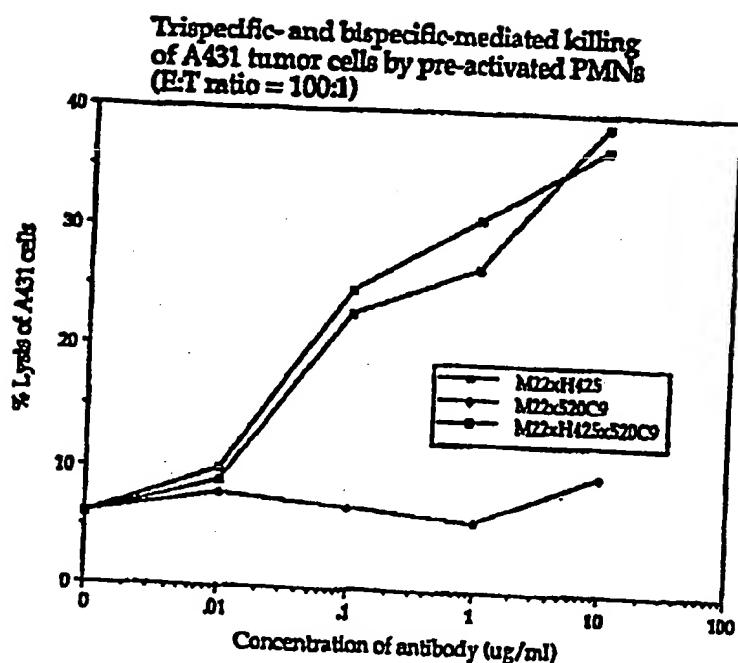


FIGURE 22

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## WHOLE BLOOD MODULATION ASSAY

Collect Whole Blood in ACD Vacutainer Tubes

Add 1 ml of Whole Blood/Well to Teflon Inserts in 24-Well Plates

22 x 22 x 32

Incubate for 4 Hours at 37°C, 5% CO<sub>2</sub>Harvest Blood from Each Well and Lyse RBCs With KHC0<sub>3</sub> Lysing Buffer

Wash Buffycoat Cells 3 Times and Transfer to 96 - Well U-Bottom Plates

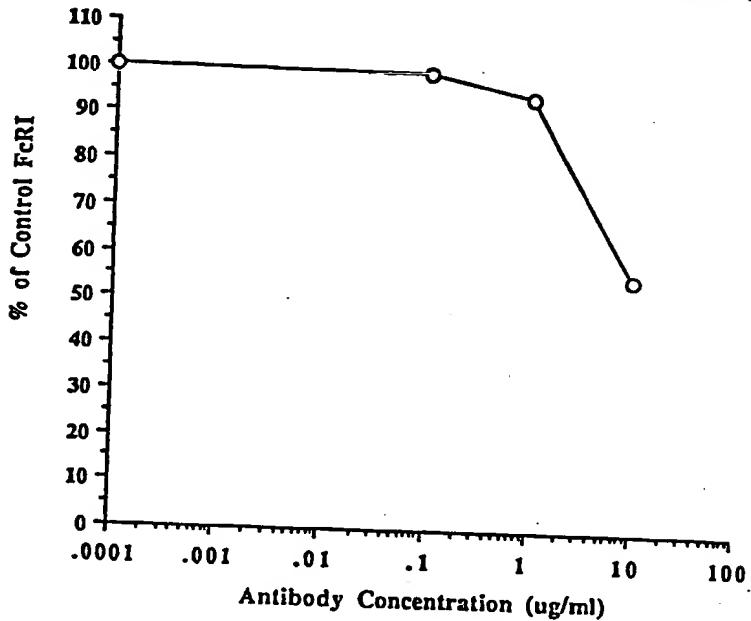
Add 100 µl of goat anti-Human IgG-PE/Well

Incubate for 1 Hour at 4°C (with shaking)

Fix Cells with 2% Paraformaldehyde

Test Samples Using Flow Cytometer

FIGURE 23A

**MODULATION ASSAY USING m22xm22x32 TRISPECIFIC ANTIBOD:****FIGURE 23B**

**A. Synthetic Oligonucleotides coding wildtype(TrT830) and mutant(TrT833S) TrT peptides:**

TrT830: QVIKANNSKRYGTEL ( *εεεα υρ νο: 6* )  
 5'-TCG AGC CAG TAC ATC AAG GCA ATC TCC AAG TTC ATC AGC AAG CTC TGA-3' ( *εεεα υρ νο: 7* )  
 3'-CG GTC ATC TAC TAC TTC CCG TTA AGC TTC ATC AAG TAC CCG TAC TGA CTC AAG ATC CCA-5' ( *εεεα υρ νο: 8* )  
 TrT833S: QVYNSNSKRYGTEL ( *εεεα υρ νο: 9* )  
 5'-TCG AGC CAG TAC ATC AGC GCG AAT TCC AAG TTC ATC CCG AAG CTC TGA-3' ( *εεεα υρ νο: 10* )  
 3'-CG GTC ATG TAC TCC GCG TTA AGG TTC AAG CCG TAC TGA CTC AAG ATC CCG-5' ( *εεεα υρ νο: 11* )

**B. Final R22 chimeric constructs**

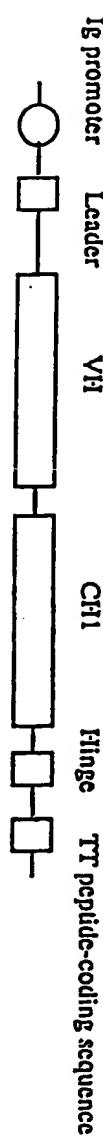
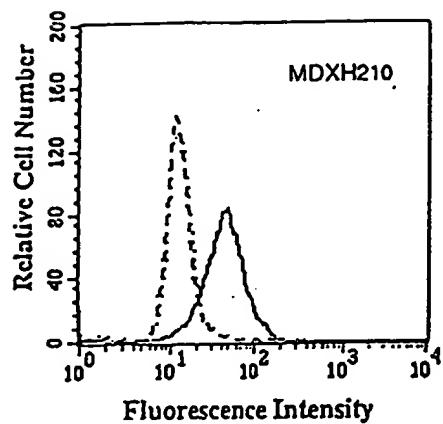
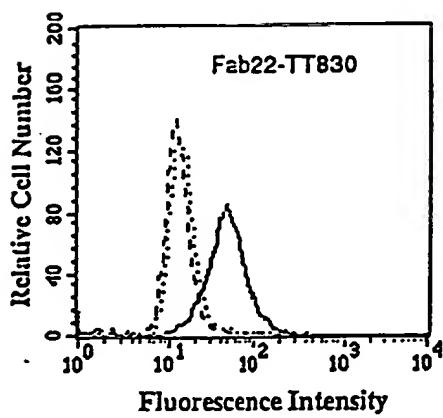


FIGURE 24

A



B



C

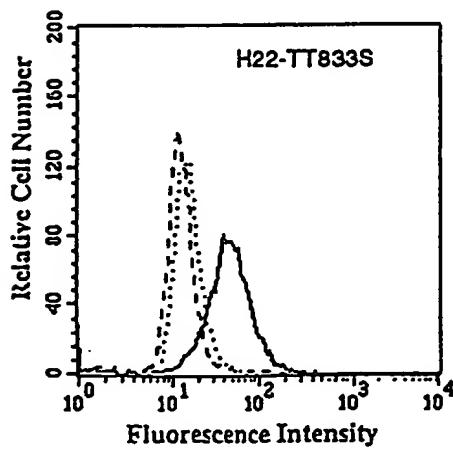


FIGURE 25

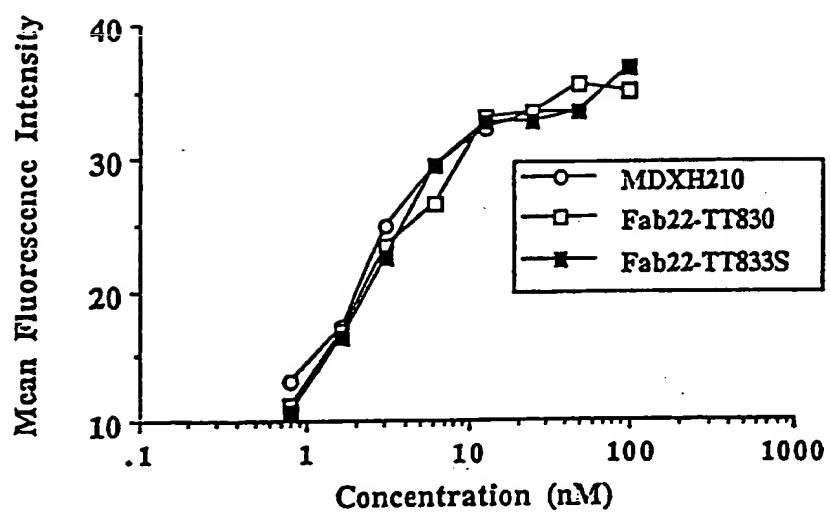


FIGURE 26

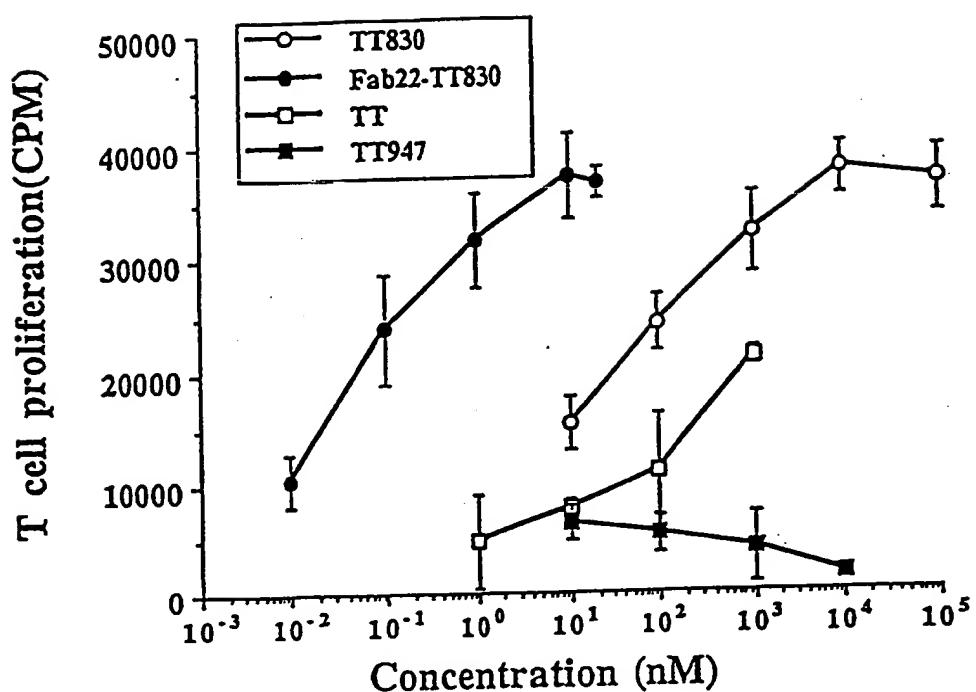


FIGURE 27

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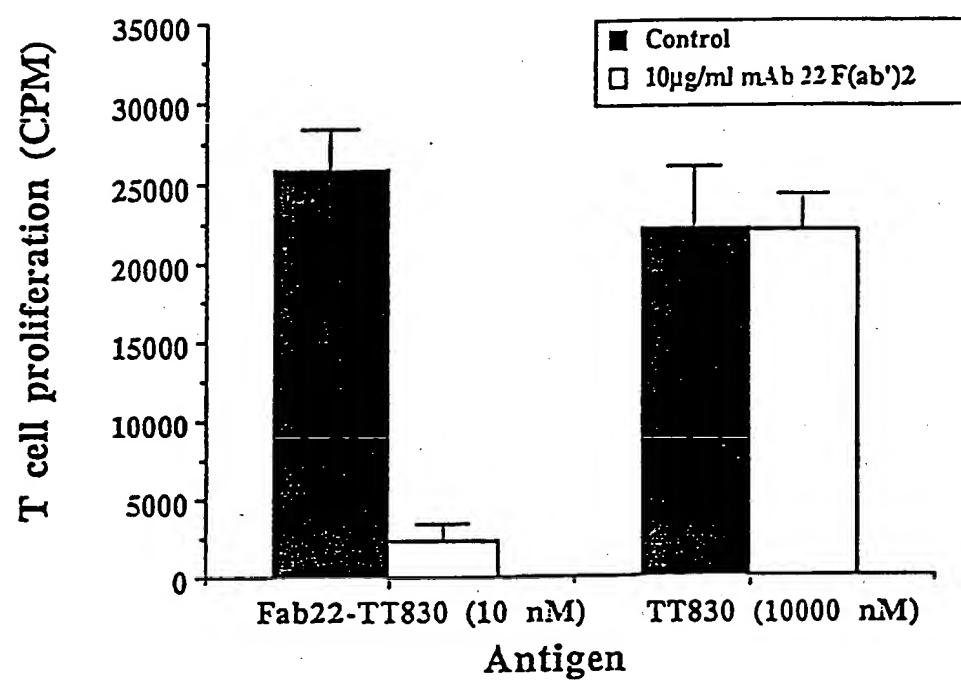


FIGURE 28

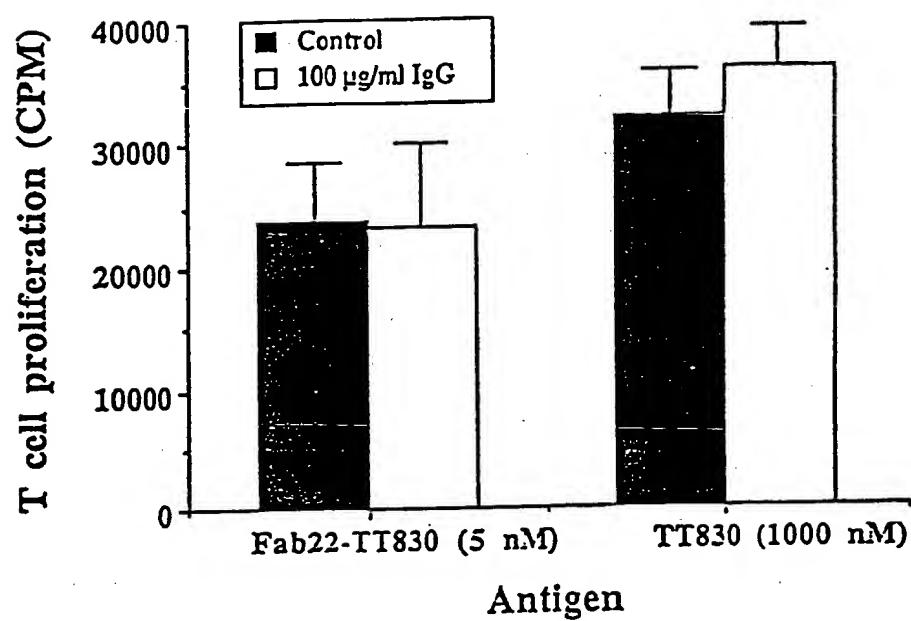


FIGURE 29

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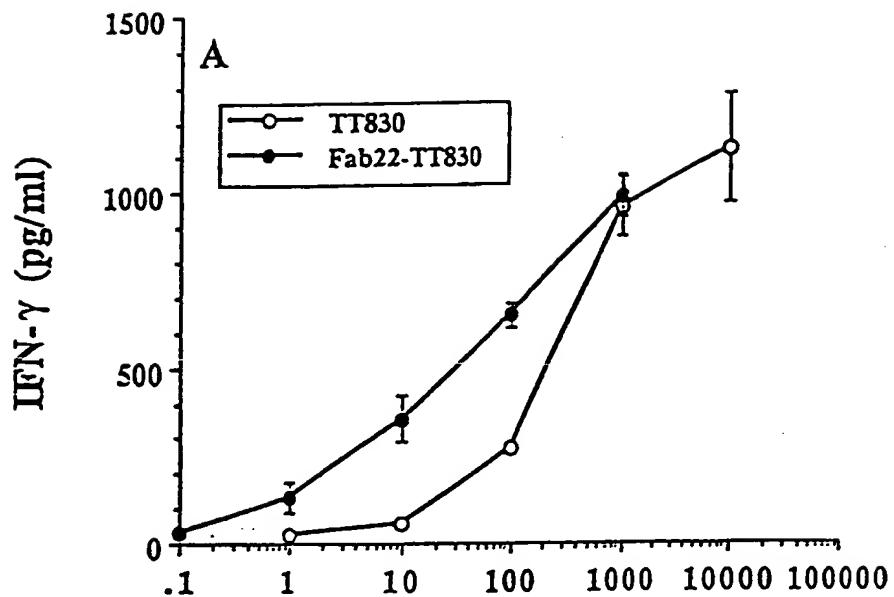


FIGURE 30A

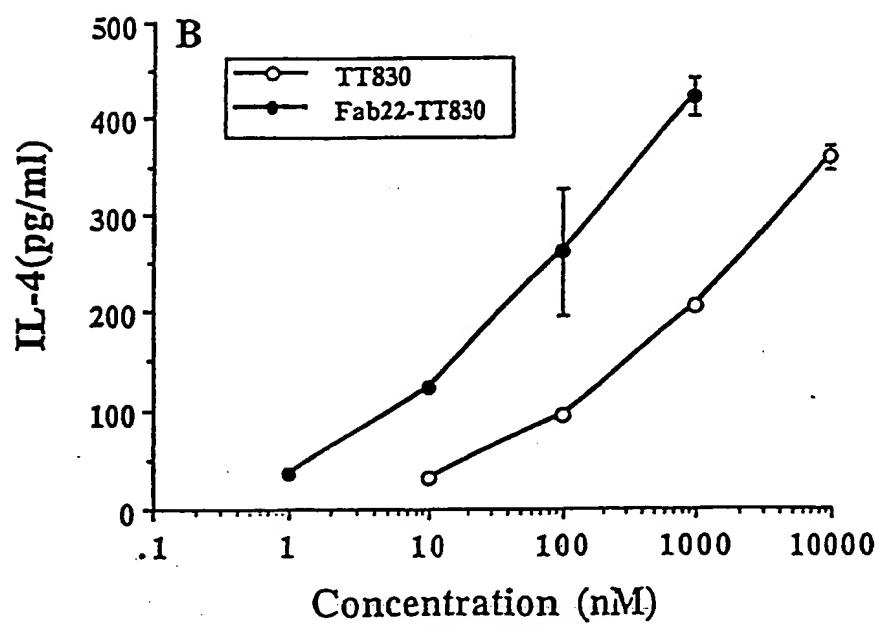


FIGURE 30B

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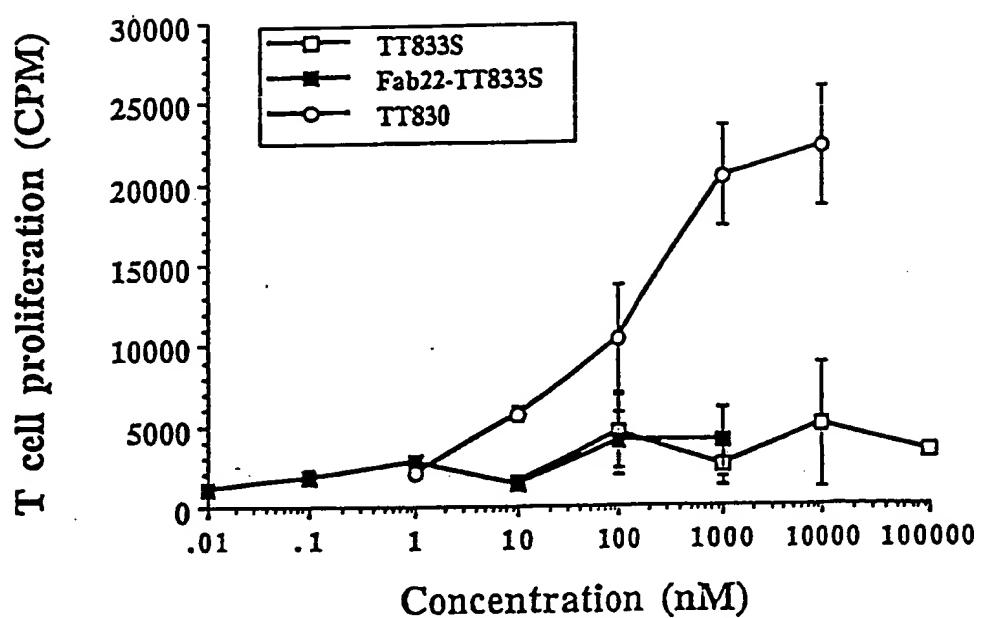


FIGURE 31

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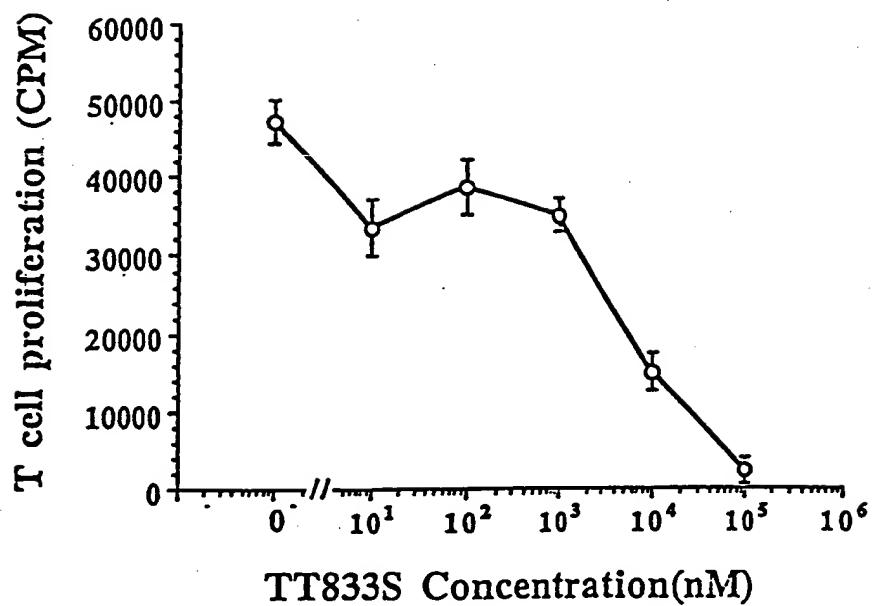


FIGURE 32

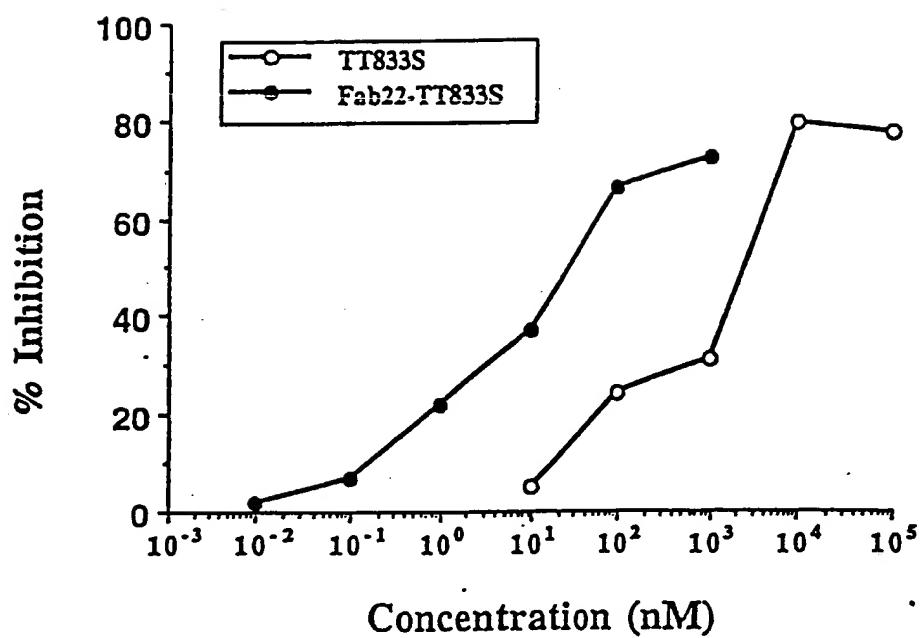


FIGURE 33

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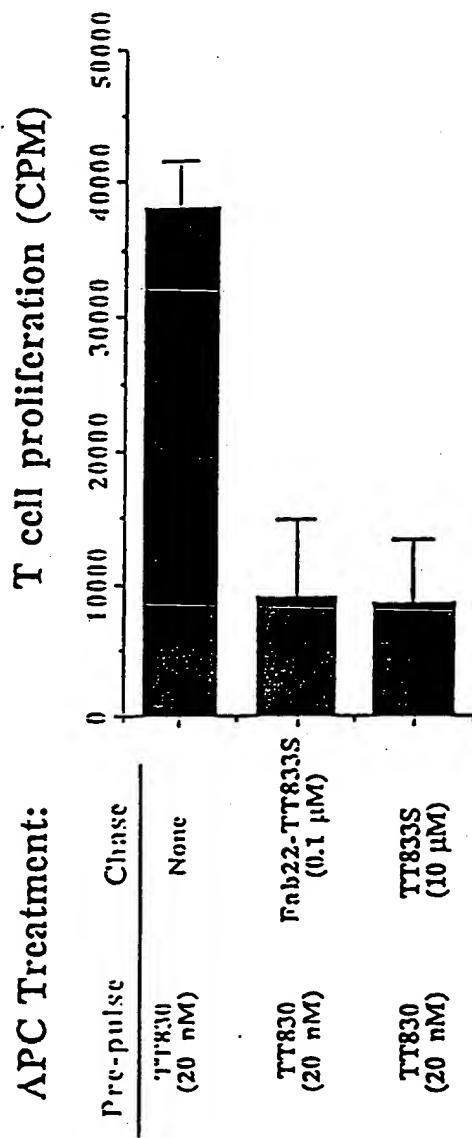


FIGURE 34

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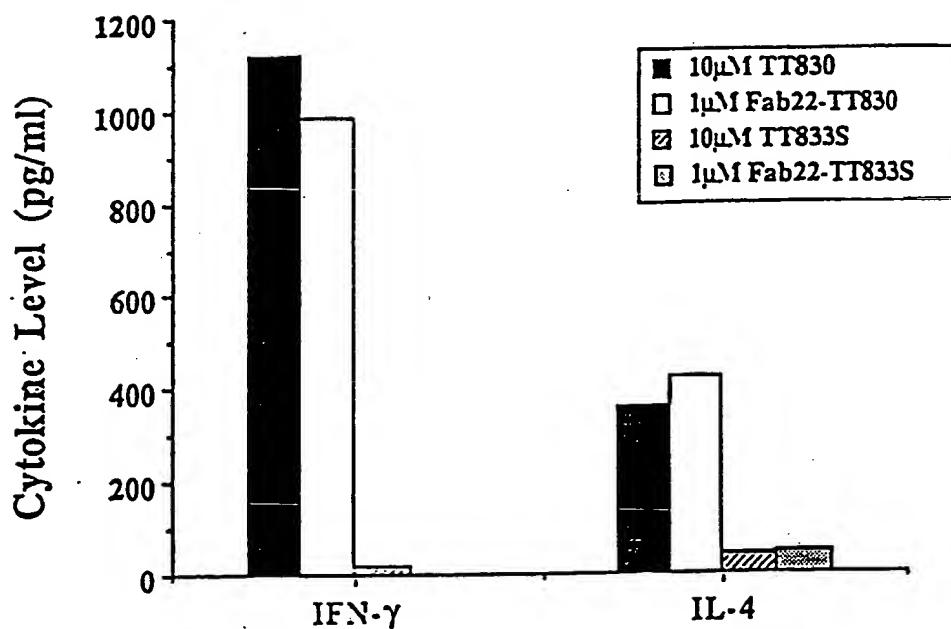


FIGURE 35

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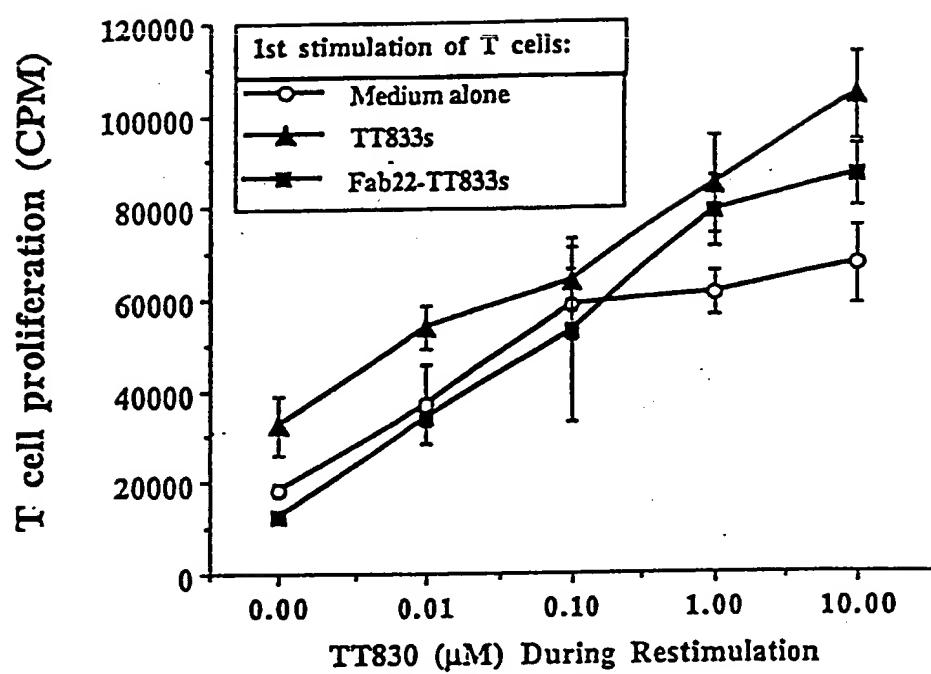
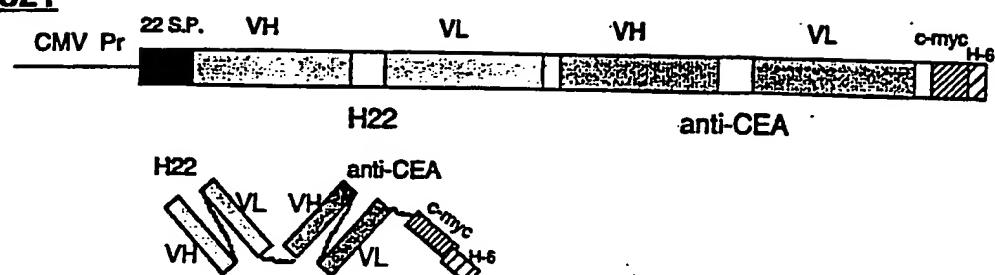
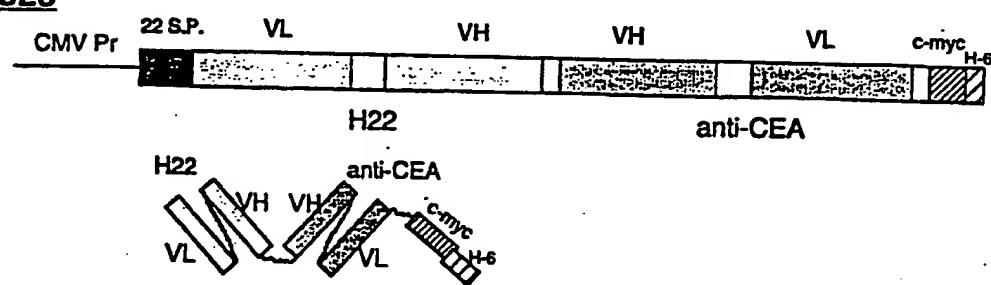
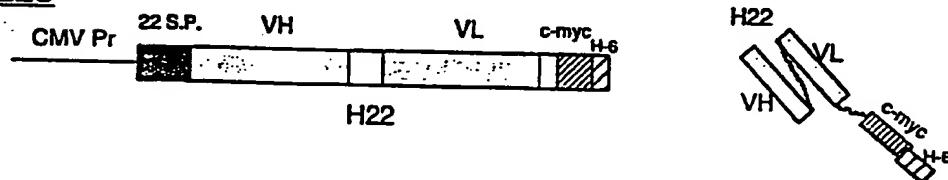


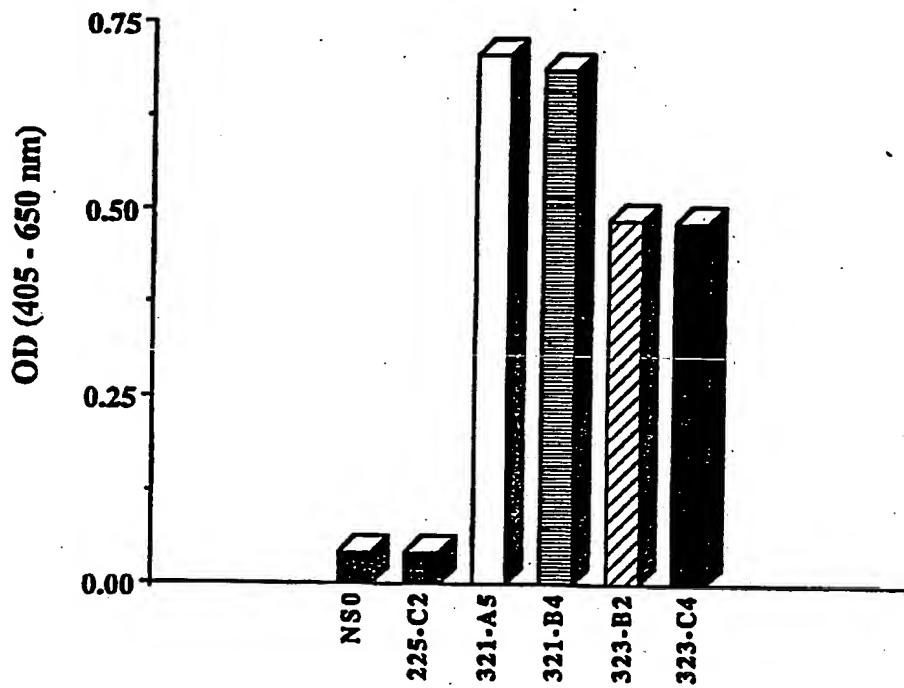
FIGURE 36

**321****323****225**

= (Gly-4, Ser)-3 linker

= (Gly-4, Ser)-1 linker

FIGURE 37

**CEA/FcRI Bispecific eLISA****FIGURE 38**

225 SEQ IDN 0:13

Sequence Range: 1 to 923

10 20 30 40 50  
 \* \* \* \* \*  
 AAGCTTCACC ATG GGA TCG AGC TGT ATC ATC CTC TTC TTG GTG GCC ACA GCT  
 TTGAAAGTGG TAC CCT ACC TCG ACA TAG TAG GAG AAG AAC CAC CGG TGT CGA  
 Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala>  
a a a TRANSLATION OF H22 SFV (225) a a a>  
 60 70 80 90 100  
 \* \* \* \* \*  
 ACC GGT GTC CAC TCC GAT ATC CAA CTG GTG GAG AGC GGT GGA GGT GTT  
 TGG CCA CAG GTG AGG CTA TAG GTT GAC CAC CTC TCG CCA CCT CCA CAA  
 Thr Gly Val His Ser (Asp Ile) Gln Leu Val Glu Ser Gly Gly Gly Val>  
a a a TRANSLATION OF H22 SFV (225) a a a>  
 110 120 130 140  
 \* \* \* \* \*  
 GTG CAA CCT GGC CGG TCC CTG CGC CTG TCC TGC TCC TCG TCT GGC TTC  
 CAC GTT GGA CCG GCC AGG GAC GCG GAC AGG AGG AGC AGA CCG AAG  
 Val Gln Pro Gly Arg Ser Leu Arg Leu Ser Cys Ser Ser Ser Gly Phe>  
a a a TRANSLATION OF H22 SFV (225) a a a>  
 150 160 170 180 190  
 \* \* \* \* \*  
 ATT TTC AGT GAC AAT TAC ATG TAT TGG GTG AGA CAG GCA CCT GGA AAA  
 TAA AAG TCA CTG TTA ATG TAC ATA ACC CAC TCT GTC CGT GGA CCT TTT  
 Ile Phe Ser Asp Asn Tyr Met Tyr Trp Val Arg Gln Ala Pro Gly Lys>  
a a a TRANSLATION OF H22 SFV (225) a a a>  
 200 210 220 230 240  
 \* \* \* \* \*  
 GGT CTT GAG TGG GTT GCA ACC ATT AGT GAT GGT GGT AGT TAC ACC TAC  
 CCA GAA CTC ACC CAA CGT TGG TAA TCA CTA CCA CCA TCA ATG TGG ATG  
 Gly Leu Glu Trp Val Ala Thr Ile Ser Asp Gly Gly Ser Tyr Thr Tyr>  
a a a TRANSLATION OF H22 SFV (225) a a a>  
 250 260 270 280 290  
 \* \* \* \* \*  
 TAT CCA GAC AGT GTG AAG GGA AGA TTT ACA ATA TCG AGA GAC AAC AGC  
 ATA GGT CTG TCA CAC TTC CCT TCT AAA TGT TAT AGC TCT CTG TTG TCG  
 Tyr Pro Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser>  
a a a TRANSLATION OF H22 SFV (225) a a a>  
 300 310 320 330 340  
 \* \* \* \* \*  
 AAG AAC ACA TTG TTC CTG CAA ATG GAC ACC CTG AGA CCC GAA GAC ACC  
 TTC TTG TGT AAC AAG GAC GTT TAC CTG TCG GAC TCT GGG CTT CTG TGG  
 Lys Asn Thr Leu Phe Leu Gln Met Asp Ser Leu Arg Pro Glu Asp Thr>  
a a a TRANSLATION OF H22 SFV (225) a a a>  
 350 360 370 380  
 \* \* \* \* \*  
 GGG GTC TAT TTT TGT GCA AGA GGC TAC TAT AGG TAC GAG GGG GGT ATG  
 CCC CAG ATA AAA ACA CGT TCT CCG ATG ATA TCC ATG CTC CCC CGA TAC  
 Gly Val Tyr Phe Cys Ala Arg Gly Tyr Tyr Arg Tyr Glu Gly Ala Met>  
a a a TRANSLATION OF H22 SFV (225) a a a>  
 390 400 410 420 430  
 \* \* \* \* \*  
 GAC TAC TGG GGC CAA GGG ACC CCG GTC ACC GTG AGC TCA GGA GGT GGC  
 CTG ATG ACC CCG GTT CCC TGG GGC CAG TGG CAC TCG AGT CCT CCA CGG  
H22V1 Linker

FIGURE 39A

Asp Tyr Trp Gly Gln Gly Thr Pro Val Thr Val Ser Ser Gly Gly Gly  
 a a a a TRANSLATION OF H22 SFV (225) a a a  
 440 450 460 470 480  
 GGC TCC GGA GGT GGA GGC AGC GGA GGG GGC GGA TCC GAC ATC CAG CTG  
 CCC AGG CCT CCA CCT CCG TCG CCT CCC CCG CCT AGG CTG TAG GTC GAC  
 Gly Ser Gly Gly Gly Ser Gly Gly Ser Asp Ile Gln Leu  
 a a a a TRANSLATION OF H22 SFV (225) a a a a  
 490 500 510 520 530  
 ACC CAG AGC CCA AGC AGC CTG AGC GCC AGC GTG GGT GAC AGA GTG ACC  
 TGG GTC TCG GGT TCG TCG GAC TCG CGG TCG CAC CCA CTG TCT CAC TGG  
 Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp Arg Val Thr  
 a a a a TRANSLATION OF H22 SFV (225) a a a a  
 540 550 560 570 580  
 ATC ACC TGT AAG TCC AGT CAA AGT GTT TTA TAC AGT TCA AAT CAG AAG  
 TAG TGG ACA TTC AGG TCA GTT TCA CAA AAT ATG TCA AGT TTA GTC TTC  
 Ile Thr Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser Ser Asn Gln Lys  
 a a a a TRANSLATION OF H22 SFV (225) a a a a  
 590 600 610 620  
 AAC TAC TTG GCC TGG TAC CAG CAG AAG CCA GGT AAG GCT CCA AAG CTG  
 TTG ATG AAC CGG ACC ATG GTC GTC TTC GGT CCA TTC CGA CGT TTC GAC  
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 630 640 650 660 670  
 CTG ATC TAC TGG GCA TCC ACT AGG GAA TCT GGT GTG CCA AGC AGA TTC  
 GAC TAG ATG ACC CGT AGG TGA TCC CTT AGA CCA CAC GGT TCG TCT AAG  
 Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val Pro Ser Arg Phe  
 a a a a TRANSLATION OF H22 SFV (225) a a a a  
 680 690 700 710 720  
 AGC GGT AGC GGT ACC GGT ACC GAC TTC ACC TTC ACC ATC AGC AGC CTC  
 TCG CCA TCG CCA TCG CGG CTG AAG TGG AAG TGG TAG TCG TCG GAG  
 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Leu  
 a a a a TRANSLATION OF H22 SFV (225) a a a a  
 730 740 750 760 770  
 CAG CCA GAG GAC ATC GCC ACC TAC TAC TGG CAT CAA TAC CTC TCC TCG  
 GTC GGT CTC CTG TAG CGG TGG ATG ATG AGC GTA GTT ATG GAG AGC AGC  
 Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys His Gln Tyr Leu Ser Ser  
 a a a a TRANSLATION OF H22 SFV (225) a a a a  
 780 790 800 810 820  
 TGG ACG TTC GGC CAA GGG ACC AAG GTG GAA ATC AAA TCT AGC TGC TCG  
 ACC TGC AAG CCG GTT CCC TGG TTC CAC CTT TAG TTT AGA TCG AGC AGC  
 Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Ser Ser Cys Ser  
 a a a a TRANSLATION OF H22 SFV (225) a a a a  
 830 840 850 860  
 AGC GGA GGC GGG GGT AGC GAT ATC GCG GCC GCA GAA CAG AAA CTC ATC  
 a a a a TRANSLATION OF H22 SFV (225) a a a a

FIGURE 39B

Linker + MCS → C-myc tag  
 TCG CCT CCG CCC CCA TCG CTA TAG CGC CGG CGT CTT GTC TTT GAG TAG  
 Ser Gly Gly Gly Ser Asp Ile Ala Ala Ala Glu Gln Lys Leu Ile>  
 \_\_\_\_\_ a a a a TRANSLATION OF H22 SFV (225) a a a a

920  
\* \* \*  
TGATTCT AGA  
ACTAAAGA TCT

**FIGURE 39C**

Sequence Range: 1 to 1679

321 SEQ ID NO: 15

10 20 30 40 50

AAGCTTCACC ATG GGA TGG AGC TGT ATC ATC CTC TTC TGG GTG GCC ACA GCT  
 TTCGAAGTGG TAC CCT ACC TCG ACA TAG TAG GAG AAG AAC CAC CGG TGT CGA  
 Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala>  
a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a >

60 <sup>5' end</sup> 70 <sup>70</sup> H22 VH 80 90 100

ACC GGT GTC CAC TCC GAT ATC CAA CTG GTG GAG AGC GGT GGA GGT GTT  
 TGG CCA CAG GTG AGG CTA TAG GTT GAC CAC CTC TCG CCA CCT CCA CAA  
 Thr Gly Val His Ser Asp Ile Gln Leu Val Glu Ser Gly Gly Gly Val>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a >

110 120 130 140

GTG CAA CCT GGC CGG TCC CTG CGC CTG TCC TGC TCC TCG TCT GGC TTC  
 CAC GTT GGA CCG GCC AGG GAC GCG GAC AGG ACG AGG AGC AGA CCG AAG  
 Val Gln Pro Gly Arg Ser Leu Arg Leu Ser Cys Ser Ser Ser Gly Phe>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a >

150 160 170 180 190

ATT TTC ACT GAC AAT TAC ATG TAT TGG GTG AGA CAG GCA CCT GGA AAA  
 TAA AAG TCA CTG TTA AAG TAC ATA ACC CAC TCT GTC CGT GGA CCT TTT  
 Ile Phe Ser Asp Asn Tyr Met Tyr Trp Val Arg Gln Ala Pro Gly Lys>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a >

200 210 220 230 240

GGT CTT GAG TGG GTT GCA ACC ATT AGT GAT GGT GGT AGT TAC ACC TAC  
 CCA GAA CTC ACC CAA CGT TGG TAA TCA CTA CCA CCA TCA ATG TGG ATG  
 Gly Leu Glu Trp Val Ala Thr Ile Ser Asp Gly Gly Ser Tyr Thr Tyr>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a >

250 260 270 280 290

TAT CCA GAC AGT GTG AAG GGA AGA TTT ACA ATA TCG AGA GAC AAC AGC  
 ATA GGT CTG TCA CAC TTC CCT TCT AAA TGT TAT AGC TCT CTG TTG TCG  
 Tyr Pro Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a >

300 310 320 330 340

AAG AAC ACA TTG TTC CTG CAA ATG GAC AGC CTG AGA CCC GAA GAC ACC  
 TTC TTG TGT AAC AAG GAC GTT TAC CTG TCG GAC TCT GGG CTT CTG TGG  
 Lys Asn Thr Leu Phe Leu Gln Met Asp Ser Leu Arg Pro Glu Asp Thr>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a >

350 360 370 380

GGG GTC TAT TTT TGT GCA AGA GGC TAC TAT AGG TAC GAG GGG GCT ATG  
 CCC CAG ATA AAA ACA CGT TCT CCG ATG ATA TCC ATG CTC CCC CGA TAC  
 Gly Val Tyr Phe Cys Ala Arg Gly Tyr Tyr Arg Tyr Glu Gly Ala Met>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a >

390 400 410 420 H22 <sup>VH</sup> 430 L14ken

GAC TAC TGG GGC CAA GGG ACC CCG GTC ACC GTG AGC TCA GGA CGT GGC  
 CTG ATG ACC CCG GTT CCC TGG GGC CAG TGG CAC TCG AGT CCT CCA CGG

FIGURE 40A

321

H22 VH

Linker

Asp Tyr Trp Gly Gln Gly Thr Pro Val Thr Val Ser Ser Gly Gly Gly  
a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

440

450

460

Linker

480

H22 VL

GGC TCC GGA GGT GGA GGC AGC GGA GGG GGC GGA TCC GAC ATC CAG CTG  
 CGG AGG CCT CCA CCT CCC TCG CCT CCC CCC CCT AGG CTG TAG GTC GAC  
 Gly Ser Gly Gly Ser Gly Gly Gly Ser Asp Ile Gln Leu  
a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

490

500

510

520

530

ACC CAG AGC CCA AGC AGC CTG AGC GCC AGC GTG GGT GAC AGA GTG ACC  
 TGG GTC TCG GGT TCG TCG GAC TCG CGG TCG CAC CCA CTG TCT CAC TGG  
 Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp Arg Val Thr  
a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

540

550

560

570

580

ATC ACC TGT AAG TCC AGT CAA AGT GTT TTA TAC AGT TCA AAT CAG AAG  
 TAG TGG ACA TTC AGG TCA GTT TCA CAA AAT ATG TCA AGT TTA GTC TTC  
 Ile Thr Cys Lys Ser Ser Gln Ser Val Leu Tyr Ser Ser Asn Gln Lys  
a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

590

600

610

620

AAC TAC TTG GCC TCG TAC CAG CAG AAG CCA GGT AAG GCT CCA AAG CTG  
 TTG ATG AAC CGG ACC ATG GTC GTC TTC GGT CCA TTC CGA GGT TTC GAC  
 Asn Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu  
a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

630

640

650

660

670

CTG ATC TAC TGG GCA TCC ACT AGG GAA TCT GGT GTG CCA ACC AGA TTC  
 GAC TAG ATG ACC CGT AGG TGA TCC CTT AGA CCA CAC GGT TCG TCT AAG  
 Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val Pro Ser Arg Phe  
a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

680

690

700

710

720

AGC GGT AGC GGT AGC GGT ACC GAC TTC ACC TTC ACC ATC AGC AGC CTC  
 TCG CCA TCG CCA TCG CCA TGG CTG AAC TCG AAC TGG TAG TCG TCG GAG  
 Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Leu  
a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

730

740

750

760

770

CAG CCA GAG GAC ATC GCC ACC TAC TAC TGC CAT CAA TAC CTC TCC TCG  
 GTC GGT CTC CTG TAG CGG TGG ATG ATG ACG GTC GTT ATG GAG AGG AGC  
 Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys His Gln Tyr Leu Ser Ser  
a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

780

790

800

H22

Linker

820

TGG ACG TTC GGC CAA GGG ACC AAG GTG GAA ATC AAA TCT AGC TGC TCG  
 ACC TGC AAG CGG GTT CCC TGG TTC CAC CTT TAG TTT AGA TCG AGC AGC  
 Trp Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Ser Ser Cys Ser  
a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

830

Linker

840

MFE-23

VL

Linker

850

860

AGC CGA GGC GGG GGT AGC GAT ATC AAA CTG CAG CAG TCT GGG GCA GAA

(321)

L<sub>4</sub>ker → MFE-23 - VH

TCG CCT CCG CCC CCA TCG CTA TAG TTT GAC GTC GTC AGA CCC CGT CTT  
 Ser Gly Gly Gly Ser Asp Ile Lys Leu Gln Gln Ser Gly Ala Glu  
 → a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a →

870 880 Gln 890 Val. 900 910

CTT GTG AGG TCA GGG ACC TCA GTC AAG TTG TCC TGC ACA GCT TCT GGC  
 GAA CAC TCC AGT CCC TGG AGT CAG TTC AAC AGG ACG TGT CGA AGA CCG  
 Leu Val Arg Ser Gly Thr Ser Val Lys Leu Ser Cys Thr Ala Ser Gly  
 → a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a →

920 930 940 950 960

TTC AAC ATT AAA GAC TCC TAT ATG CAC TGG TTG AGG CAG GGG CCT GAA  
 AAG TTG TAA TTT CTG AGG ATA TAC GTG ACC AAC TCC GTC CCC GGA CTT  
 Phe Asn Ile Lys Asp Ser Tyr Met His Trp Leu Arg Gln Gly Pro Glu  
 → a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a →

970 980 990 1000 1010

CAG GGC CTG GAG TGG ATT GGA TGG ATT GAT CCT GAG AAT GGT GAT ACT  
 GTC CCG GAC CTC ACC TAA CCT ACC TAA CTA GGA CTC TTA CCA CTA TGA  
 Gln Gly Leu Glu Trp Ile Gly Trp Ile Asp Pro Glu Asn Gly Asp Thr  
 → a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a →

1020 1030 1040 1050 1060

GAA TAT GCC CCG AAG TTC CAG GGC AAG GCC ACT TTT ACT ACA GAC ACA  
 CTT ATA CGG GGC TTC AAG GTC CCC TTC CCG TGA AAA TGA TGT CTG TGT  
 Glu Tyr Ala Pro Lys Phe Gln Gly Lys Ala Thr Phe Thr Thr Asp Thr  
 → a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a →

1070 1080 1090 1100

TCC TCC AAC ACA GCC TAC CTG CAG CTG ACC AGC CTG ACA TCT GAG GAC  
 AGG AGG TTG TGT CCG ATG GAC GTC GAC TCG TCG GAC TGT AGA CTC CTG  
 Ser Ser Asn Thr Ala Tyr Leu Gln Leu Ser Ser Leu Thr Ser Glu Asp  
 → a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a →

1110 1120 1130 1140 1150

ACT GCC GTC TAT TAT TGT AAT GAG GGG ACT CCC ACT GGG CGG TAC TAC  
 TGA CGG CAG ATA ATA ACA TTA CTC CCC TGA GGC TGA CCC GGC ATG ATG  
 Thr Ala Val Tyr Tyr Cys Asn Glu Gly Thr Pro Thr Gly Pro Tyr Tyr  
 → a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a →

1160 1170 1180 1190 MFE-23 1200

TTT GAC TAC TGG GGC CAA GGG ACC AGC GTC ACC GTC TCC TCA GGT GGA  
 AAA CTG ATG ACC CCG GTT CCC TGG TGC CAG TGG CAG AGT CCA CCT  
 Phe Asp Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Gly  
 → a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a →

1210 1220 1230 1240 1250

GGC GGT TCA GGC GGA GGT GGC TCT GGC GGT GGC GGA TCA GAA AAT GTG VL  
 CCG CCA AGT CGG CCT CCA CCG AGA CCG CCA CCG CCT AGT CTT TTA CAC  
 Gly Gly Ser Gly Gly Ser Gly Gly Gly Ser Glu Asn Val  
 → a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a →

1260 1270 1280 1290 1300

(321)

CTC ACC CAG TCT CCA GCA ATC ATG TCT GCA TCT CCA GGG GAG AAG GTC  
 GAG TGG GTC AGA GGT CGT TAG TAC AGA CGT AGA GGT CCC CTC TTC CAG  
 Leu Thr Gln Ser Pro Ala Ile Met Ser Ala Ser Pro Gly Glu Lys Val>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

1310                    1320                    1330                    1340  
 \*                    \*                    \*                    \*  
 ACC ATA ACC TCC AGT GCC AGC TCA AGT GTC AGT TAC ATG CAC TGG TTC  
 TGG TAT TGG ACG TCA CGG TCG AGT TCA CAT TCA ATG TAC GTG ACC AAC  
 Thr Ile Thr Cys Ser Ala Ser Ser Val Ser Tyr Met His Trp Phe>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

1350                    1360                    1370                    1380                    1390  
 \*                    \*                    \*                    \*                    \*  
 CAG CAG AAG CCA GGC ACT TCT CCC AAA CTC TGG ATT TAT AGC ACA TCC  
 GTC GTC TTC GGT CGG TGA AGA GGG TTT GAG ACC TAA ATA TCG TGT AGG  
 Gln Gln Lys Pro Gly Thr Ser Pro Lys Leu Trp Ile Tyr Ser Thr Ser>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

1400                    1410                    1420                    1430                    1440  
 \*                    \*                    \*                    \*                    \*  
 AAC CTG GCT TCT GGA GTC CCT GCT CGC TTC AGT GGC AGT GGA TCT GGG  
 TTG GAC CGA AGA CCT CAG GGA CGA CGG AAG TCA CGG TCA CCT AGA CCC  
 Asn Leu Ala Ser Gly Val Pro Ala Arg Phe Ser Gly Ser Gly Ser Gly>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

1450                    1460                    1470                    1480                    1490  
 \*                    \*                    \*                    \*                    \*  
 ACC TCT TAC TCT CTC ACA ATC AGC CGA ATG GAG GCT GAA GAT GCT GCC  
 TGG AGA ATG AGA GAG TGT TAG TCG GCT TAC CTC CGA CTT CTA CGA CGG  
 Thr Ser Tyr Ser Leu Thr Ile Ser Arg Met Glu Ala Glu Asp Ala Ala>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

1500                    1510                    1520                    1530                    1540  
 \*                    \*                    \*                    \*                    \*  
 ACT TAT TAC TGC CAG CAA AGG AGT AGT TAC CCA CTC ACG TTC GGT GCT  
 TGA ATA ATG AGC GTC GTT TCC TCA TCA ATG GGT GAG TGC AAG CCA CGA  
 Thr Tyr Tyr Cys Gln Gln Arg Ser Ser Tyr Pro Leu Thr Phe Gly Ala>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

1550                    1560                    1570                    1580  
 \*                    \*                    \*                    \*  
 GGC ACC AAG CTG GAG CTG AAA CGG GCG GCA GGC TCG AGC GGA GGC CGC  
 CCG TGG TTC GAC CTC GAC TTT GCC CGC CGT CCG AGC TCG CCT CCG CCC  
 Gly Thr Lys Leu Glu Leu Lys Arg Ala Ala Gly Ser Ser Gly Gly Gly>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

1590                    1600                    1610                    1620                    1630  
 \*                    \*                    \*                    \*                    \*  
 GGT AGC GAT ATC GCG GCC GCA GAA CAG AAA CTC ATC TCA GAA GAG GAT  
 CCA TCG CTA TAG CGC CGG CGT CTT GTC TTT GAG TAG AGT CTT CTC CTA  
 Gly Ser Asp Ile Ala Ala Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp>  
a a TRANSLATION OF H22/MFE-23 SC-BSAB (321) [A] a a >

1640                    1650                    1660                    1670  
 C-<sup>14</sup> C-<sup>14</sup>            \*                    \*                    \*  
 CTG AAT GGC GCC GCA CAT CAC CAT CAT CAC CAT T GATTCTAGA  
 GAC TTA CGG CGG CGT GTA GTG GTA GTG GTA A CTAAGATCT  
 Leu Asn Gly Ala Ala His His His His His>  
TRANSLATION OF H22/MFE-23 SC-BSAB (3) >

## INTERNATIONAL SEARCH REPORT

International Application No  
PCT/US 96/09988

A. CLASSIFICATION OF SUBJECT MATTER  
IPC 6 C07K16/46 A61K39/395 //C07K16/28, C07K16/32, C07K14/485,  
C07K16/30

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)  
IPC 6 C07K A61K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	<p>WO 94 10332 A (MEDAREX) 11 May 1994 cited in the application</p> <p>see the whole document</p> <p>---</p> <p>-/-</p>	1-10, 15, 17-20, 23-25, 33-35

Further documents are listed in the continuation of box C.

Patent family members are listed in annex.

## \* Special categories of cited documents :

- \*A\* document defining the general state of the art which is not considered to be of particular relevance
- \*E\* earlier document but published on or after the international filing date
- \*L\* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- \*O\* document referring to an oral disclosure, use, exhibition or other means
- \*P\* document published prior to the international filing date but later than the priority date claimed

\*T\* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

\*X\* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

\*Y\* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.

\*Z\* document member of the same patent family

1 Date of the actual completion of the international search

17 October 1996

Date of mailing of the international search report

05.11.96

## Name and mailing address of the ISA

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Fax (+ 31-70) 340-3016

## Authorized officer

Nooij, F

## INTERNATIONAL SEARCH REPORT

Inte onal Application No  
PCT/US 96/09988

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	IMMUNOMETHODS, vol. 4, no. 1, February 1994, SAN DIEGO, CA, USA, pages 72-81, XP000608179 M. FANGER ET AL.: "Production and use of anti-FcR bispecific antibodies." see page 73, left-hand column, line 15 - right-hand column, line 4 see page 76, right-hand column, line 9 - line 21 see page 76, right-hand column, line 43 - line 51 see page 77, left-hand column, line 40 - line 56 see page 78, right-hand column, line 8 - page 79, left-hand column, line 9 see page 79, right-hand column, line 9 - line 27 see figure 1 ---	1-9, 11-20, 22-25, 32-35
X	IMMUNOBIOLOGY, vol. 191, no. 2-3, September 1994, STUTTGART, GERMANY, pages 250-251, XP000608202 R. REPP ET AL.: "G-CSF stimulated neutrophils as effector cells in immunotherapy with a bispecific antibody to FcgammaRI and to HER-2/neu (MDX210): preclinical studies." see the whole document ---	1,15,17, 18, 23-25,33
X	WO 92 05793 A (MEDAREX) 16 April 1992  see examples see claims ---	1,15, 23-25,35
X	WO 91 05871 A (MEDAREX) 2 May 1991  see examples see claims ---	1,15,17, 23-25, 32,33
X	JOURNAL OF GENERAL VIROLOGY, vol. 75, no. 6, June 1994, CAMBRIDGE, GB, pages 1451-1456, XP002016265 A. MABONDZO ET AL.: "Antibody-dependent cellular cytotoxicity and neutralization of human immunodeficiency virus type 1 by high affinity cross-linking of gp41 to human macrophage Fc IgG receptor using bispecific antibody." see abstract ---	1-7,9, 15,20, 22-25,35
1		-/-

## INTERNATIONAL SEARCH REPORT

International Application No  
PCT/US 96/09988

## C(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	<p>CLINICAL CANCER RESEARCH, vol. 1, no. 4, April 1995, USA, pages 425-434, XP000608205</p> <p>J. CHEN ET AL.: "An immunoconjugate of Lys3-bombesin and monoclonal antibody 22 can specifically induce Fc<sub>gamma</sub>RI (CD64)-dependent monocyte- and neutrophil-mediated lysis of small cell carcinoma of the lung cells."</p> <p>see abstract see discussion</p> <p>---</p>	1,15,17, 18, 23-25,33
A	<p>EP 0 255 249 A (TRUSTEES OF DARTMOUTH COLLEGE) 3 February 1988</p> <p>cited in the application</p> <p>see exemplification</p> <p>see claims</p> <p>---</p>	1-35
P,X	<p>THE JOURNAL OF IMMUNOLOGY, vol. 155, no. 10, 15 November 1995, BALTIMORE, MD, USA, pages 4996-5002, XP002016266</p> <p>R. GRAZIANO ET AL.: "Construction and characterization of a humanized anti-gamma-Ig receptor type I (Fc<sub>gamma</sub>RI) monoclonal antibody."</p> <p>see abstract see page 5001, right-hand column, line 11 - line 35</p> <p>-----</p>	1-9, 11-15, 17-20, 22-25, 33,35

## INTERNATIONAL SEARCH REPORT

International application No.

PCT/US 96/09988

## Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1.  Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:

**Remark:** Although claim(s) 33 - 35 is(are) directed to a method of treatment of the human/animal body, the search has been carried out and based on the alleged effects of the compound/composition.

2.  Claims Nos.: because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:

3.  Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

## Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1.  As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.

2.  As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.

3.  As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:

4.  No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

## Remark on Protest

The additional search fees were accompanied by the applicant's protest.

No protest accompanied the payment of additional search fees.

## INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/US 96/09988

Patent document cited in search report	Publication date	Patent family member(s)		Publication date
WO-A-9410332	11-05-94	AU-A-	5542894	24-05-94
		CA-A-	2148578	11-05-94
		EP-A-	0668924	30-08-95
WO-A-9205793	16-04-92	AU-B-	667460	28-03-96
		AU-A-	8869491	28-04-92
		CA-A-	2093022	06-04-92
		EP-A-	0553244	04-08-93
		JP-T-	6502410	17-03-94
WO-A-9105871	02-05-91	CA-A-	2069960	21-04-91
		EP-A-	0595798	11-05-94
EP-A-255249	03-02-88	US-A-	4954617	04-09-90
		AT-T-	120802	15-04-95
		AU-B-	695771	24-01-91
		AU-A-	7527187	14-01-88
		CA-A-	1319899	06-07-93
		DE-D-	3751214	11-05-95
		DE-T-	3751214	16-11-95
		EP-A-	0629703	21-12-94
		ES-T-	2072851	01-08-95
		IL-A-	101475	31-07-94
		WO-A-	8800052	14-01-88
		JP-T-	1500195	26-01-89